GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: September 11, 2005, 18:03:37 ; Search time 1720.68 Seconds
(without alignments)
9546.401 Million cell updates/sec

Title: US-09-403-107-145_COPY_1_339
Perfect score: 339
Sequence: 1 gaggtgcagctgctcgagtc.....ccctggtcaccgtctcctca 339

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 4708233 seqs, 24227607955 residues

Total number of hits satisfying chosen parameters: 9416466

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:*
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2: gb_htg:*
3: gb_in:*
4: gb_om:*
5: gb_ov:*
6: gb_pat:*
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12: gb_sy:*
13: gb_un:*
14: gb_vi:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	339	100.0	372	6	A84378	A84378 Sequence 14
2	339	100.0	372	6	BD075295	BD075295 Novel met
3	304.4	89.8	339	9	AJ627239	AJ627239 Homo sapi
4	300.6	88.7	339	9	AF035024	AF035024 Homo sapi
5	292.6	86.3	362	9	HSA308463	AJ308463 Homo sapi
6	291.8	86.1	345	9	AB063888	AB063888 Homo sapi
7	291.6	86.0	342	9	HSIGHVCA	Z31686 H.sapiens (
8	291.6	86.0	437	9	HSE5435	Z14203 H.sapiens r
9	286.6	84.5	342	9	HSA579107	AJ579107 Homo sapi
10	285.6	84.2	398	9	HSE55310	Z14168 H.sapiens r
11	285.4	84.2	345	9	AB067097	AB067097 Homo sapi
12	285.2	84.1	336	9	HSA579114	AJ579114 Homo sapi
13	285.2	84.1	354	9	HSA300793	AJ300793 Homo sapi
14	284.4	83.9	357	6	AX300024	AX300024 Sequence
15	284	83.8	376	9	AF115111	AF115111 Homo sapi
16	283.8	83.7	354	9	AF062288	AF062288 Homo sapi
17	283.4	83.6	399	9	HSIGHXX17	X65899 H.sapiens m
18	283	83.5	363	9	AF062116	AF062116 Homo sapi
19	282.6	83.4	354	9	HSA245291	AJ245291 Homo sapi
20	282.6	83.4	414	9	AF062277	AF062277 Homo sapi
21	282.4	83.3	351	9	HSA245286	AJ245286 Homo sapi
22	282.4	83.3	351	9	HSA245307	AJ245307 Homo sapi
23	282.4	83.3	351	9	HSA579139	AJ579139 Homo sapi
24	282.4	83.3	351	9	HSA579151	AJ579151 Homo sapi
25	282.2	83.2	345	9	AB063830	AB063830 Homo sapi
26	282	83.2	392	9	HSE5535	Z14192 H.sapiens r
27	281.8	83.1	351	9	AY530317	AY530317 Homo sapi
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31	281.4	83.0	360	9	HSA245351	AJ245351 Homo sapi
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33	281.2	82.9	357	9	AJ627243	AJ627243 Homo sapi
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36	280.8	82.8	357	9	AF062287	AF062287 Homo sapi
37	280.4	82.7	348	9	AF035021	AF035021 Homo sapi
38	279.6	82.5	357	9	HSA579099	AJ579099 Homo sapi
39	279.4	82.4	425	9	AY392931	AY392931 Homo sapi
40	279	82.3	348	9	HSA245279	AJ245279 Homo sapi
41	278.4	82.1	381	6	BD139667	BD139667 A novel m
42	278.4	82.1	381	6	AX003759	AX003759 Sequence
43	278.4	82.1	414	6	A84376	A84376 Sequence 14
44	278.4	82.1	414	6	BD075294	BD075294 Novel met
45	278.4	82.1	1630	6	BD222938	BD222938 Heteromin

ALIGNMENTS

RESULT 1
A84378
LOCUS A84378 372 bp DNA linear PAT 21-JAN-2000
DEFINITION Sequence 145 from Patent WO9846645.
ACCESSION A84378
VERSION A84378.1 GI:6733301
KEYWORDS .
SOURCE unidentified
ORGANISM unidentified
unclassified.
REFERENCE 1 (bases 1 to 372)
AUTHORS Kufer,P. and Raum,T.
TITLE NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN RECEPTORS AND USES THEREOF
JOURNAL Patent: WO 9846645-A 145 22-OCT-1998;
KUFER PETER (DE); RAUM TOBIAS (DE)
FEATURES Location/Qualifiers
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/db_xref="taxon:32644"
CDS <1. .>372
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/protein_id="CAB69289.1"
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/translation="EVQLLESGGVVVQPGGSLRLSCAASGFTFDDYAMHWVRQAPGKG
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ORIGN

Query Match 100.0%; Score 339; DB 6; Length 372;
Best Local Similarity 100.0%; Pred. No. 9.2e-100;
Matches 339; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
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Db          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
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Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
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Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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RESULT 2
BD075295
LOCUS       BD075295                372 bp    DNA     linear   PAT 27-AUG-2002
DEFINITION  Novel method for the production of anti-human antigen receptors and
            uses thereof.
ACCESSION   BD075295
VERSION     BD075295.1  GI:22620898
KEYWORDS    JP 2001519824-A/24.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 372)
  AUTHORS   Kufer,P. and Raum,T.
  TITLE     Novel method for the production of anti-human antigen receptors and
            uses thereof
  JOURNAL   Patent: JP 2001519824-A 24 23-OCT-2001;
            MICROMET AG
COMMENT     OS   Homo sapiens (human)
            PN   JP 2001519824-A/24
            PD   23-OCT-2001
            PF   14-APR-1998 JP 1998543494
            PR   14-APR-1997 EP   97106109.8
            PI   PETER KUFER,TOBIAS RAUM
            PC   C07K16/00,C07K16/30,A61K39/395
            CC   Novel method for the production of anti-human antigen  CC
                  receptors and uses
            CC   thereof
            FH   Key             Location/Qualifiers
            FT   CDS             (1). .(372).
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                      /mol_type="genomic DNA"
                      /db_xref="taxon:9606"
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  Best Local Similarity   100.0%;  Pred. No. 9.2e-100;
  Matches  339;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
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Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
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Db        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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Db        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
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Db        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300

Qy        301 GGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
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Db        301 GGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339


RESULT 3
AJ627239
LOCUS       AJ627239                339 bp    mRNA    linear   PRI 30-JUN-2004
DEFINITION  Homo sapiens partial mRNA for IgD immunoglobulin heavy chain
            variable region (IGHV3-30 gene), clone T3.3.13.
ACCESSION   AJ627239
VERSION     AJ627239.1  GI:49523851
KEYWORDS    IGHV3-30 gene; immunoglobulin heavy chain; variable region.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Dono,M.
  TITLE     Characterization of a novel CD5+ B cell population
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 339)
  AUTHORS   Dono,M.
  TITLE     Direct Submission
  JOURNAL   Submitted (10-FEB-2004) Dono M., Oncologia Medica C, Istituto
            Nazionale Ricerca sul Cancro, L.go R. Benzi 10, Italy, 16132, ITALY
FEATURES              Location/Qualifiers
     source           1. .339
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                      /cell_type="B cell"
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                      /gene="IGHV3-30"
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  Best Local Similarity   93.8%;  Pred. No. 2.1e-88;
  Matches  317;  Conservative    0;  Mismatches   21;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
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Db        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAACATCCTG 301

Qy        302 GCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
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Db        302 AGACCTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339


RESULT 4
AF035024
LOCUS       AF035024                339 bp    mRNA    linear   PRI 24-SEP-1999
DEFINITION  Homo sapiens clone MCE11H myosin-reactive immunoglobulin heavy
            chain variable region mRNA, partial cds.
ACCESSION   AF035024
VERSION     AF035024.1  GI:5921600
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 339)
  AUTHORS   Wu,X., Liu,B., Van der Merwe,P.L., Kalis,N.N., Berney,S.M. and
            Young,D.C.
  TITLE     Myosin-reactive autoantibodies in rheumatic carditis and normal
            fetus
  JOURNAL   Clin. Immunol. Immunopathol. 87 (2), 184-192 (1998)
  MEDLINE   98277139
   PUBMED   9614934
REFERENCE   2  (bases 1 to 339)
  AUTHORS   Young,D.C.
  TITLE     Direct Submission
  JOURNAL   Submitted (19-NOV-1997) Department of Pathology and Laboratory
            Medicine, University of Texas Health Science Center, 6431 Fannin,
            Houston, TX 77030, USA
FEATURES             Location/Qualifiers
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ORIGIN

  Query Match           88.7%;  Score 300.6;  DB 9;  Length 339;
  Best Local Similarity   92.9%;  Pred. No. 3.6e-87;
  Matches  315;  Conservative    0;  Mismatches   24;  Indels    0;  Gaps    0;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
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Db          1 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGGGGTCCCTGAGACTC 60

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
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Db        121 CCAGGCAAGGGGCTGGAGTGGGTGGCATTTATACGGTATGATGGAAGTAATAAATACTAT 180

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
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RESULT 5
HSA308463
LOCUS       HSA308463               362 bp    mRNA    linear   PRI 06-FEB-2002
DEFINITION  Homo sapiens partial mRNA for immunoglobulin heavy chain variable
            region (IGHV3 gene), clone TEG4.
ACCESSION   AJ308463
VERSION     AJ308463.1  GI:12734095
KEYWORDS    IGHV3 gene; immunoglobulin heavy chain; variable region.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Jacobin,M.J., Laroche-Traineau,J., Little,M., Keller,A., Peter,K.,
            Welschof,M., Nurden,A. and Clofent-Sanchez,G.
  TITLE     Human IgG monoclonal anti-alpha(IIb)beta(3)-binding fragments
            derived from immunized donors using phage display
  JOURNAL   J. Immunol. 168 (4), 2035-2045 (2002)
  MEDLINE   21681719
   PUBMED   11823541
REFERENCE   2  (bases 1 to 362)
  AUTHORS   Clofent-Sanchez,G.
  TITLE     Direct Submission
  JOURNAL   Submitted (06-FEB-2001) Clofent-Sanchez G., UMR5533, Cnrs, Hopital
            Cardiologique, Av de Magellan, 33604 Pessac, FRANCE
FEATURES             Location/Qualifiers
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                     /note="alphaIIbbeta3 integrin specific antibody"
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ORIGIN

  Query Match           86.3%;  Score 292.6;  DB 9;  Length 362;
  Best Local Similarity   92.8%;  Pred. No. 1.5e-84;
  Matches  320;  Conservative    0;  Mismatches   19;  Indels    6;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
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Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
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Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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Db        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAGAAAGTCT 300

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Db        301 TACTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 345
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RESULT 6
AB063888
LOCUS AB063888 345 bp mRNA linear PRI 02-JUL-2002
DEFINITION Homo sapiens IGH mRNA for immunoglobulin heavy chain VHDJ region, partial cds, clone:H238.
ACCESSION AB063888
VERSION AB063888.1 GI:21668977
KEYWORDS .
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1
AUTHORS Akahori,Y., Iba,Y., Morino,K., Shinohara,M., Hirono,Y., Kakita,M., Suzuki,K., Torii,H., Ukai,Y., Honda,T., Katsumi,H., Okada,J., Miura,K. and Kurosawa,Y.
TITLE Construction and characterization of antibody libraries: isolation of therapeutic human antibodies and application to functional genomics
JOURNAL Unpublished
REFERENCE 2 (bases 1 to 345)
AUTHORS Kurosawa,Y.
TITLE Direct Submission
JOURNAL Submitted (25-JUN-2001) Yoshikazu Kurosawa, Institute for Comprehensive Medical Science, Fujita Health University; Kutsukake-cho, Toyoake 470-1192, Japan
(E-mail:kurosawa@fujita-hu.ac.jp, Tel:81-562-93-9387)
COMMENT Please visit our web site
URL:http://www.fujita-hu.ac.jp/immunity/.
FEATURES Location/Qualifiers
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="H238"
/clone_lib="AIMS4"
/note="mixture of tissues:tonsils, umbilical cords, peripheral blood and bone marrow"
gene 1. .345
/gene="IGH"
CDS <1. .>345
/gene="IGH"
/codon_start=1
/product="immunoglobulin heavy chain VHDJ region"
/protein_id="BAC01516.1"
/db_xref="GI:21668978"
/translation="EVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKG
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DYWGQGTLVTVSS"
ORIGIN

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  Query Match            86.1%;  Score 291.8;  DB 9;  Length 345;
  Best Local Similarity  93.3%;  Pred. No. 2.8e-84;
  Matches  318;  Conservative    0;  Mismatches   17;  Indels    6;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
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Db          1 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 60

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
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Db         61 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 120

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
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Db        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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Db        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAA------ 294
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGGGCGG 300

Qy        295 AAGGAAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
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Db        301 ACTGGGGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTC 341
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RESULT 7
HSIGHVCA
LOCUS HSIGHVCA 342 bp DNA linear PRI 05-AUG-1999
DEFINITION H.sapiens (VHIL) gene for immunoglobulin heavy chain variable region.
ACCESSION Z31686
VERSION Z31686.1 GI:509782
KEYWORDS immunoglobulin heavy chain; immunoglobulin heavy chain variable region.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 342)
AUTHORS Figini,M., Marks,J.D., Winter,G. and Griffiths,A.D.
TITLE In vitro assembly of repertoires of antibody chains on the surface of phage by renaturation
JOURNAL J. Mol. Biol. 239 (1), 68-78 (1994)
MEDLINE 94254092
PUBMED 8196048
REFERENCE 2 (bases 1 to 342)
AUTHORS Griffiths,A.D.
TITLE Direct Submission
JOURNAL Submitted (30-MAR-1994) A.D. Griffiths, MRC Centre for Protein Engineering, Hills Road, Cambridge, CB2 2QH, U.K
FEATURES Location/Qualifiers
source 1. .342
/organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
/cell_type="peripheral lymphocyte"
/tissue_type="blood"
/dev_stage="adult"
CDS 1. .342
/note="Protein sequence is in conflict with the conceptual translation"
/codon_start=1
/product="immunoglobulin heavy chain chain variable region (VH-IL)"
/protein_id="CAA83491.1"
/db_xref="GI:1335143"
/translation="EVQLVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKG
LEWVAVISYDGSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDWGD
YWGQGTLVTVSS"

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ORIGIN

  Query Match             86.0%;  Score 291.6;  DB 9;  Length 342;
  Best Local Similarity   93.5%;  Pred. No. 3.3e-84;
  Matches  316;  Conservative    0;  Mismatches   19;  Indels    3;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
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Db          1 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 60

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
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Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
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Db        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCAATAAATACTAT 180

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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Db        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA---G 297
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Db        241 CTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGACTGG 300

Qy        298 GAAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
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Db        301 GGAGACTACTGGGGCCAGGGAACCCTGGTCACCGTGTC 338


RESULT 8
HSE5435
LOCUS       HSE5435                  437 bp    mRNA    linear   PRI 15-MAR-1993
DEFINITION  H.sapiens rearranged Ig heavy chain variable region (VDJ).
ACCESSION   Z14203 X65741
VERSION     Z14203.1  GI:30965
KEYWORDS    Ig D-segment; Ig heavy chain; Ig J-segment; Ig V-segment; Ig
            variable region; immunoglobulin.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 437)
  AUTHORS   Cuisinier,A.M., Gauthier,L., Boubli,L., Fougereau,M. and
            Tonnelle,C.
  TITLE     Mechanisms that generate human immunoglobulin diversity operate
            from the 8th week of gestation in fetal liver
  JOURNAL   Eur. J. Immunol. 23 (1), 110-118 (1993)
  MEDLINE   93122076
   PUBMED   8419161
REFERENCE   2  (bases 1 to 437)
  AUTHORS   Tonnelle,C.
  TITLE     Direct Submission
  JOURNAL   Submitted (09-JUN-1992) C. Tonnelle, Centre d'Immunologie
            Marseille-Luminy, Case 906, 13288 Marseille Cedex 9, FRANCE
FEATURES             Location/Qualifiers
     source          1. .437
                     /organism="Homo sapiens"
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                     /tissue_type="feral liver"
                     /clone_lib="E54"
     CDS             36. .>437
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                     /protein_id="CAA78572.1"
                     /db_xref="GI:30966"
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                     /note="Ig D-segment"
     misc_feature    399. .437
                     /note="Ig J-segment"
ORIGIN

  Query Match             86.0%;  Score 291.6;  DB 9;  Length 437;
  Best Local Similarity   92.7%;  Pred. No. 3.4e-84;
  Matches  319;  Conservative    0;  Mismatches   19;  Indels    6;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Db         94 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 153

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Db        154 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGCTATGCACTGGGTCCGCCAGGCTC 213

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        214 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCAATAAATACTACG 273

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Db        274 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 333

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAG---- 297
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Db        334 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAGAGAGCC 393

Qy        298 --GAAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
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Db        394 GAGGGGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 437


RESULT 9
HSA579107
LOCUS       HSA579107                342 bp    mRNA    linear   PRI 01-JUN-2004
DEFINITION  Homo sapiens partial mRNA for IgM immunoglobulin heavy chain
            variable region (IGHV3-33 gene), clone T1.3.14.
ACCESSION   AJ579107
VERSION     AJ579107.1  GI:47846392
KEYWORDS    IGHV3-33 gene; immunoglobulin heavy chain; variable region.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Dono,M.
  TITLE     Characterization of a novel CD5+ B cell population
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 342)
  AUTHORS   Dono,M.
  TITLE     Direct Submission
  JOURNAL   Submitted (07-MAY-2003) Dono M., Oncologia Medica C., Istituto
            Nazionale Ricerca sul Cancro, L. go R. Benzi 10, Genova, 16132,
            ITALY
FEATURES             Location/Qualifiers
     source          1. .342
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                     /db_xref="taxon:9606"
                     /chromosome="14"
                     /clone="T1.3.14"
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     gene            1. .342
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  Query Match             84.5%;  Score 286.6;  DB 9;  Length 342;
  Best Local Similarity   92.1%;  Pred. No. 1.4e-82;
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Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Db         62 CCTGTGCAGCGTCTGGATTCACCTTTAGTAACTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATGGTATGATGGAAGTAATGCATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Db        182 CAGACTCCGTGAACGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACATGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA---GG 298
              |||||||||||||||||||||| |||||||||||||||||||||||||||||  | |   ||
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGCTACATCGGG 301

Qy        299 AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
              | | | |||||||||||||||||||||||||||||||||||
Db        302 AGGCCAACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 342


RESULT 10
HSE55310
LOCUS       HSE55310                398 bp    mRNA    linear   PRI 15-MAR-1993
DEFINITION  H.sapiens rearranged Ig heavy chain variable region (VDJ).
ACCESSION   Z14168 X65741
VERSION     Z14168.1  GI:30999
KEYWORDS    Ig D-segment; Ig heavy chain; Ig J-segment; Ig V-segment; Ig
            variable region; immunoglobulin.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 398)
  AUTHORS   Cuisinier,A.M., Gauthier,L., Boubli,L., Fougereau,M. and
            Tonnelle,C.
  TITLE     Mechanisms that generate human immunoglobulin diversity operate
            from the 8th week of gestation in fetal liver
  JOURNAL   Eur. J. Immunol. 23 (1), 110-118 (1993)
  MEDLINE   93122076
   PUBMED   8419161
REFERENCE   2  (bases 1 to 398)
  AUTHORS   Tonnelle,C.
  TITLE     Direct Submission
  JOURNAL   Submitted (09-JUN-1992) C. Tonnelle, Centre d'Immunologie
            Marseille-Luminy, Case 906, 13288 Marseille Cedex 9, FRANCE
FEATURES             Location/Qualifiers
     source          1. .398
                     /organism="Homo sapiens"
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                     /db_xref="taxon:9606"
                     /chromosome="14"
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                     /clone_lib="E55"
     CDS             <1. .>398
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                     /db_xref="GI:31000"
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                     AEDTAVYYCAKDLFYYFDYWGQGTLVTVSS"
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                     /note="signal-peptide site"
     misc_feature    48. .342
                     /note="Ig V-segment"
     misc_feature    343. .351
                     /note="Ig D-segment"
     misc_feature    352. .398
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ORIGIN

  Query Match             84.2%;  Score 285.6;  DB 9;  Length 398;
  Best Local Similarity   91.1%;  Pred. No. 3.1e-82;
  Matches  319;  Conservative    0;  Mismatches   19;  Indels   12;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Db         49 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 108

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |   |||| ||||||||||||||||||||||||
Db        109 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 168

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        169 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 228

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        229 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 288

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAA------- 294
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Db        289 TGCAAATGAACGGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGATCTCT 348

Qy        295 -----AAGGAAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                  |    | |||||||||||||||||||||||||||||||||||||
Db        349 TCTACTACTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 398


RESULT 11
AB067097
LOCUS       AB067097                345 bp    mRNA    linear   PRI 02-JUL-2002
DEFINITION  Homo sapiens IGH mRNA for immunoglobulin heavy chain VHDJ region,
            partial cds, clone:aims0184h.
ACCESSION   AB067097
VERSION     AB067097.1  GI:21670250
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Akahori,Y., Iba,Y., Morino,K., Shinohara,M., Hirono,Y., Kakita,M.,
            Suzuki,K., Torii,H., Ukai,Y., Honda,T., Katsumi,H., Okada,J.,
            Miura,K. and Kurosawa,Y.
  TITLE     Construction and characterization of antibody libraries: isolation
            of therapeutic human antibodies and application to functional
            genomics
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 345)
  AUTHORS   Kurosawa,Y.
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  TITLE     Direct Submission
  JOURNAL   Submitted (25-JUL-2001) Yoshikazu Kurosawa, Institute for
            Comprehensive Medical Science, Fujita Health University,
            Immunology; Kutsukake-cho, Toyoake, Aichi 470-1192, Japan
            (E-mail:kurosawa@fujita-hu.ac.jp, Tel:81-562-93-9387)
COMMENT     Please visit our web site
            URL:http://www.fujita-hu.ac.jp/immunity/.
FEATURES             Location/Qualifiers
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                     /db_xref="taxon:9606"
                     /clone="aims0184h"
                     /clone_lib="AIMS4"
                     /note="mixture of tissues:tonsils, umbilical cords,
                     peripheral blood and bone marrow"
     gene            1. .345
                     /gene="IGH"
     CDS             <1. .>345
                     /gene="IGH"
                     /codon_start=1
                     /product="immunoglobulin heavy chain VHDJ region"
                     /protein_id="BAC02135.1"
                     /db_xref="GI:21670251"
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                     LEWVAVISYDGSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCALGGNF
                     NYWGQGTLVTVSS"
ORIGIN

  Query Match             84.2%;  Score 285.4;  DB 9;  Length 345;
  Best Local Similarity   92.1%;  Pred. No. 3.5e-82;
  Matches  314;  Conservative    0;  Mismatches   21;  Indels    6;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              ||||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||
Db          1 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 60

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              |||||||||||||||||||||||||    |   |||| ||||||||||||||||||||||
Db         61 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 120

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCG------AAA 294
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Db        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGCTCGGTGGT 300

Qy        295 AAGGAAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
              ||      |||||||||||||||||||||||||||||||||
Db        301 AACTTTAACTACTGGGGCCAGGGAACCCTGGTCACCGTCTC 341


RESULT 12
HSA579114
LOCUS       HSA579114               336 bp    mRNA    linear   PRI 01-JUN-2004
DEFINITION  Homo sapiens partial mRNA for IgM immunoglobulin heavy chain
            variable region (IGHV3-30 gene), clone T1.3.23.
ACCESSION   AJ579114
VERSION     AJ579114.1  GI:47846406
KEYWORDS    IGHV3-30 gene; immunoglobulin heavy chain; variable region.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Dono,M.
  TITLE     Characterization of a novel CD5+ B cell population
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 336)
  AUTHORS   Dono,M.
  TITLE     Direct Submission
  JOURNAL   Submitted (07-MAY-2003) Dono M., Oncologia Medica C., Istituto
            Nazionale Ricerca sul Cancro, L. go R. Benzi 10, Genova, 16132,
            ITALY
FEATURES             Location/Qualifiers
     source          1. .336
                     /organism="Homo sapiens"
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ORIGIN

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  Best Local Similarity   92.3%;  Pred. No. 4.1e-82;
  Matches  312;  Conservative    0;  Mismatches   23;  Indels    3;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGAGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
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Qy        302 GCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
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RESULT 13
HSA300793
LOCUS       HSA300793               354 bp    mRNA    linear   PRI 10-APR-2001
DEFINITION  Homo sapiens partial mRNA for immunoglobulin gamma heavy chain
            variable region (IGHV3-30), clone M2-45 (m2hie45), Kawasaki disease
            patient.
ACCESSION   AJ300793
VERSION     AJ300793.1  GI:12733989
KEYWORDS    IGHV3-30 gene; immunoglobulin gamma heavy chain; kawasaki disease;
            variable region.
SOURCE      Homo sapiens (human)
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  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 354)
  AUTHORS   Leucht,S., Uttenreuther-Fischer,M.M., Gaedicke,G. and Fischer,P.
  TITLE     The B cell superantigen-like interaction of intravenous
            immunoglobin (IVIG) with Fab fragments of V(H) 3-23 and 3-30/3-30.5
            germline gene origin cloned from a patient with Kawasaki disease is
            enhanced after IVIG therapy
  JOURNAL   Clin. Immunol. 99 (1), 18-29 (2001)
  MEDLINE   21185274
   PUBMED   11286538
REFERENCE   2  (bases 1 to 354)
  AUTHORS   Fischer,P.
  TITLE     Direct Submission
  JOURNAL   Submitted (17-JAN-2001) Fischer P., Charite Children's Hospital,
            Molecular Biology Laboratory, Humboldt-University, Ziegelstr. 5-9,
            Berlin, 10117, GERMANY
FEATURES             Location/Qualifiers
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ORIGIN

  Query Match             84.1%;  Score 285.2;  DB 9;  Length 354;
  Best Local Similarity   90.7%;  Pred. No. 4.1e-82;
  Matches  321;  Conservative    0;  Mismatches   18;  Indels   15;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
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Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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Qy        301 ---------------GGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
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RESULT 14
AX300024
LOCUS       AX300024                357 bp    DNA     linear   PAT 30-NOV-2001
DEFINITION  Sequence 2 from Patent WO0185797.
ACCESSION   AX300024
VERSION     AX300024.1  GI:17381484
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Rodriguez,M., Miller,D.J. and Pease,L.R.
  TITLE     Human igm antibodies with the capability of inducing remyelination,
            and diagnostic and therapeutic uses thereof particularly in the
            central nervous system
  JOURNAL   Patent: WO 0185797-A 2 15-NOV-2001;
            MAYO FOUNDATION FOR MEDICAL EDUCATION AND RESEARCH (US)
FEATURES             Location/Qualifiers
     source          1. .357
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  Matches  322;  Conservative    0;  Mismatches   16;  Indels   18;  Gaps    1;

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Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
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Qy        299 ---------------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
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RESULT 15
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LOCUS       AF115111                376 bp    mRNA    linear   PRI 04-JAN-2000
DEFINITION  Homo sapiens clone dn5 immunoglobulin heavy chain variable region
            mRNA, partial cds.
ACCESSION   AF115111
VERSION     AF115111.1  GI:4836305
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 376)
  AUTHORS   Smithson,S.L., Srivastava,N., Hutchins,W.A. and Westerink,M.A.
  TITLE     Molecular analysis of the heavy chain of antibodies that recognize
            the capsular polysaccharide of Neisseria meningitidis in hu-PBMC
            reconstituted SCID mice and in the immunized human donor
```

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  JOURNAL   Mol. Immunol. 36 (2), 113-124 (1999)
  MEDLINE   99305028
   PUBMED   10378683
REFERENCE   2  (bases 1 to 376)
  AUTHORS   Smithson,S.L., Srivastava,N., Hutchins,W.A. and Westerink,M.A.J.
  TITLE     Direct Submission
  JOURNAL   Submitted (17-DEC-1998) Medicine, Medical College of Ohio, 3000
            Arlington Ave, Toledo, OH 43614, USA
FEATURES             Location/Qualifiers
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Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Db         62 CCTGTGCAGCGTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCG---------- 291
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Qy        292 --AAAAAGGAAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
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Search completed: September 11, 2005, 21:22:00
Job time : 1725.68 secs
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: September 11, 2005, 17:46:48 ; Search time 356.464 Seconds
(without alignments)
5629.725 Million cell updates/sec

Title: US-09-403-107-145_COPY_1_339
Perfect score: 339
Sequence: 1 gaggtgcagctgctcgagtc.....ccctggtcaccgtctcctca 339

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 4390206 seqs, 2959870667 residues

Total number of hits satisfying chosen parameters: 8780412

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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12: geneseqn2004as:*
13: geneseqn2004bs:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	339	100.0	372	2	AAV68538	Aav68538 Nucleotid
2	301.2	88.8	729	12	ADN07003	Adn07003 Human EFG
3	294.8	87.0	396	10	ADC61027	Adc61027 Human ant
4	289.2	85.3	342	12	ADQ75774	Adq75774 Anti-IL-8
5	284.4	83.9	357	12	ADI26655	Adi26655 Human ant
6	284	83.8	351	12	ADI22042	Adi22042 Anti-plat
7	284	83.8	351	12	ADI22043	Adi22043 Anti-plat
8	283.8	83.7	405	10	ADC61031	Adc61031 Human ant
9	282.4	83.3	351	12	ADI22044	Adi22044 Anti-plat
10	282.4	83.3	351	12	ADI22041	Adi22041 Anti-plat
11	282	83.2	345	12	ADI22052	Adi22052 Anti-plat
12	281	82.9	354	6	ABK88453	Abk88453 Human ant
13	280.8	82.8	351	12	ADI22046	Adi22046 Anti-plat
14	280.8	82.8	351	12	ADI22045	Adi22045 Anti-plat
15	280.6	82.8	349	13	ADS84379	Ads84379 Human ant
16	280.6	82.8	349	13	ADR68521	Adr68521 Anti-EPO-
17	280.6	82.8	404	10	ADC61047	Adc61047 Human ant
18	280.6	82.8	405	10	ADC61043	Adc61043 Human ant
19	279.2	82.4	351	12	ADI22047	Adi22047 Anti-plat
20	279	82.3	405	10	ADC61039	Adc61039 Human ant
21	278.4	82.1	381	2	AAX77236	Aax77236 Human D4.
22	278.4	82.1	414	2	AAV68537	Aav68537 Nucleotid
23	278.4	82.1	1630	3	AAZ50588	Aaz50588 HD70scFv-
24	278.4	82.1	1630	3	AAZ50587	Aaz50587 HD70scFv-
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26	277.2	81.8	339	4	AAS03438	Aas03438 DNA encod
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28	277.2	81.8	2202	13	ACN41350	Acn41350 Human dia
29	276.6	81.6	369	2	AAT60380	Aat60380 Anti-TGF
30	276.2	81.5	400	6	ABL38517	Abl38517 Human col
31	275.8	81.4	405	10	ADC61051	Adc61051 Human ant
32	275.8	81.4	405	10	ADC61111	Adc61111 Human ant
33	275.6	81.3	339	4	AAS03437	Aas03437 DNA encod
34	275	81.1	345	2	AAT60369	Aat60369 Anti-TGF
35	275	81.1	350	2	AAT60370	Aat60370 Anti-TGF
36	274.6	81.0	354	6	ABA94330	Aba94330 MAb 6-2 h
37	274.6	81.0	366	12	ADP22127	Adp22127 Human ant
38	274.4	80.9	351	6	ABK88457	Abk88457 Human ant
39	274.4	80.9	376	10	ADE28450	Ade28450 Human ant
40	274.4	80.9	675	4	AAH41661	Aah41661 Human int
41	274.4	80.9	675	4	AAH30007	Aah30007 Anti-IL8
42	274.2	80.9	349	13	ADS84391	Ads84391 Human ant
43	274.2	80.9	349	13	ADR68533	Adr68533 Anti-EPO-
44	274.2	80.9	405	10	ADC61055	Adc61055 Human ant
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ALIGNMENTS

RESULT 1
AAV68538
ID AAV68538 standard; DNA; 372 BP.
XX
AC AAV68538;
XX
DT 16-FEB-1999 (first entry)
XX
DE Nucleotide sequence of human D7.2 heavy chain variable region.
XX
KW Human; D7.2 heavy chain variable region; receptor; antigen; tumour;
KW auto-immune disease; graft rejection; allergy; inflammatory disease;
KW endocrine disease; degenerative disease; ss.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT CDS 1. .372
FT /*tag= a
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XX
PN WO9846645-A2.
XX
PD 22-OCT-1998.
XX
PF 14-APR-1998; 98WO-EP002180.
XX
PR 14-APR-1997; 97EP-00106109.
XX
PA (KUFE/) KUFER P.
PA (RAUM/) RAUM T.
XX
PI Kufer P, Raum T;
XX
DR WPI; 1998-594564/50.
DR P-PSDB; AAW80816.
XX
PT Production of anti-human antigen receptors - by selecting a combination
PT of functionally rearranged VH and VL immunoglobulin chains expressed from
PT a recombinant vector.
XX
PS Claim 9; Fig 8; 84pp; English.
XX

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CC   This is the nucleotide sequence of the human D7.2 heavy chain variable
CC   region, used in the method of the invention. for providing receptors that
CC   can be used for targeting antigens in humans without being immunogenic
CC   themselves. Such receptors can be used for treating diseases such as
CC   tumours or auto-immune diseases, graft rejection after transplantation,
CC   infectious diseases by targeting cellular receptors as well as allergic,
CC   inflammatory, endocrine and degenerative diseases by targeting key
CC   molecules involved in the pathological process
XX
SQ   Sequence 372 BP; 86 A; 95 C; 110 G; 81 T; 0 U; 0 Other;

  Query Match             100.0%;  Score 339;  DB 2;  Length 372;
  Best Local Similarity   100.0%;  Pred. No. 5.6e-91;
  Matches  339;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
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Db          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
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Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
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Db        301 GGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339


RESULT 2
ADN07003
ID   ADN07003 standard; DNA; 729 BP.
XX
AC   ADN07003;
XX
DT   01-JUL-2004  (first entry)
XX
DE   Human EFGR bs-scFv antibody associated DNA.
XX
KW   Bispecific single chain; bs-scFv; cancer;
KW   epidermal growth factor receptor; EGFR; therapy; antibody; human; ds.
XX
OS   Homo sapiens.
XX
PN   US2004071696-A1.
XX
PD   15-APR-2004.
XX
PF   04-APR-2003; 2003US-00406830.
XX
PR   05-APR-2002; 2002US-0370276P.
XX
PA   (REGC ) UNIV CALIFORNIA.
PA   (FOXC-) FOX CHASE CANCER CENT.
XX
PI   Adams GP,  Horak EM,  Weiner LM,  Marks JD;
XX
DR   WPI; 2004-328525/30.
XX
PT   Novel bispecific antibody comprising first and second antibody joined to
PT   each other and having binding specificity to different epitopes of
PT   Epidermal Growth Factor Receptor protein, useful for treating cancer.
XX
PS   Claim 14; SEQ ID NO 20; 57pp; English.
XX
CC   The present invention provides bispecific single chain (bs-scFv) antibody
CC   molecules which may be used to treat various forms of cancer associated
CC   with the overexpression of the epidermal growth factor receptor (EGFR)
CC   family. The invention is useful for specifically delivering an effector
CC   molecule to a cell bearing a receptor from EGFR protein family chosen
CC   from EGFR, HER2/neu, HER3 and HER4. The invention is useful in the
CC   treatment of cancer. The present sequence is human EFGR bs-scFv antibody
CC   associated DNA used in the invention.
XX
SQ   Sequence 729 BP; 155 A; 196 C; 216 G; 162 T; 0 U; 0 Other;

  Query Match             88.8%;  Score 301.2;  DB 12;  Length 729;
  Best Local Similarity   93.2%;  Pred. No. 1.4e-79;
  Matches  315;  Conservative    0;  Mismatches   23;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||||| |||||| |||||||| ||||||||||||||||
Db          8 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 67

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |   |||| ||||||||||||||||||||||||
Db         68 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 127

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        128 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 187

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        188 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 247

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              |||||||||||||||||||||||||||||||||| |||||||||||||   |
Db        248 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGAGTATTACTGTGCGAAGTATCCTT 307

Qy        302 GCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                 |||||||||||||||||||||||||||||||||||||
Db        308 TAAACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 345


RESULT 3
ADC61027
ID   ADC61027 standard; DNA; 396 BP.
XX
AC   ADC61027;
XX
DT   18-DEC-2003  (first entry)
XX
DE   Human anti-CD45RB monoclonal antibody DNA, SEQ ID No 54.
XX
KW   monoclonal antibody; CD45RB; antigen; cell proliferation;
KW   immunosuppressive; neuroprotective; tissue rejection; organ rejection;
KW   autoimmune disease; multiple sclerosis; human; anti-CD45RB; gene; ds.
XX
OS   Homo sapiens.
XX
PN   WO2003048327-A2.
XX
PD   12-JUN-2003.
XX
PF   02-DEC-2002; 2002WO-US038540.
XX
PR   03-DEC-2001; 2001US-0337276P.
XX
PA   (ABGE-) ABGENIX INC.
XX
PI   Foltz I,  Babcook J,  Palathumpat V,  Yang X,  King CT;
XX
DR   WPI; 2003-558954/52.
DR   P-PSDB; ADC61028.
```

```
XX
PT   New anti-CD45RB monoclonal antibody, useful for treating an autoimmune
PT   disease e.g. multiple sclerosis.
XX
PS   Disclosure; SEQ ID NO 54; 121pp; English.
XX
CC   The invention relates to a novel isolated monoclonal antibody comprising
CC   a heavy chain having a sequence chosen from one of 22 fully defined
CC   sequences comprising 135-147 amino acids, given in the specification, and
CC   is specific for CD45RB antigen. The invention further relates to:
CC   inhibiting cell proliferation associated with the expression of CD45RB
CC   antigen; and treating a disease associated with the expression of a
CC   CD45RB antigen in a patient. The monoclonal antibody has the activities
CC   of immunosuppressive and neuroprotective. The monoclonal antibody is
CC   useful for treating the rejection of a mammalian cell, tissue or organ,
CC   especially an autoimmune disease in a mammal, especially a human e.g.
CC   multiple sclerosis. This polynucleotide sequence represents a DNA
CC   encoding a human protein of the anti-CD45RB monoclonal antibody of the
CC   invention.
XX
SQ   Sequence 396 BP; 84 A; 92 C; 124 G; 96 T; 0 U; 0 Other;

  Query Match             87.0%;  Score 294.8;  DB 10;  Length 396;
  Best Local Similarity   92.0%;  Pred. No. 9e-78;
  Matches  311;  Conservative    0;  Mismatches   27;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| |||||||||| ||||||||||||||||
Db         59 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 118

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| ||||||||||||||    | | |||| ||||||||||||||||||||||||
Db        119 CCTGTGCAGCGTCTGGATTCACCTTCAGTAACTATGGCATGCACTGGGTCCGCCAGGCTC 178

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||| |||||||  |||||||||||||| | || |||||
Db        179 CAGGCAAGGGGCTGGAGTGGGTGGCTGTTATATGGTATGATGGAAGTAAGAGATTCTATG 238

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        239 CAGGCTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 298

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              |||||||||||||||||||||| |||||||||||||||||||||||||| ||    ||
Db        299 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAACTACAG 358

Qy        302 GCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
               |||||||||||||||||| ||||||||||||||||
Db        359 ACTACTGGGGCCAGGGAACCATGGTCACCGTCTCCTCA 396


RESULT 4
ADQ75774
ID   ADQ75774 standard; cDNA; 342 BP.
XX
AC   ADQ75774;
XX
DT   07-OCT-2004  (first entry)
XX
DE   Anti-IL-8 antibody, VH 3-33, VH coding sequence.
XX
KW   antibody; heavy chain; light chain; variable region; constant region; VH;
KW   VL; monoclonal antibody; MAb; interleukin-8; IL-8;
KW   complementarity determining region; CDR; inflammation;
KW   hyperproliferation; skin disorder; PPP; psoriasis; plaque psoriasis;
KW   guttate type psoriasis; bullous skin disease; bullous pemphigoid;
KW   contact dermatitis; eczema; erythematosus; atopic dermatitis;
KW   immune disorder; autoimmune disorder; inflammatory disease;
KW   psoriatic arthritis; systemic scleroderma; sclerosis;
KW   inflammatory bowel disease; IBD; Crohn's disease; ulcerative colitis;
KW   acute lung injury; acute respiratory distress syndrome;
KW   adult respiratory distress syndrome; meningitis; encephalitis; uveitis;
KW   multiple myeloma; glomerulonephritis; nephritis; asthma; atherosclerosis;
KW   leukocyte adhesion deficiency; multiple sclerosis; Raynaud's syndrome;
KW   Sjogren's syndrome; juvenile onset diabetes; Reiter's disease;
KW   Behcet's disease; immune complex nephritis; IgA nephropathy;
KW   IgM polyneuropathy; immune-mediated thrombocytopenia;
KW   acute idiopathic thrombocytopenic purpura;
KW   chronic idiopathic thrombocytopenic purpura; haemolytic anaemia;
KW   myasthenia gravis; lupus nephritis; lupus erythematosus;
KW   rheumatoid arthritis; ankylosing spondylitis; pemphigus; Graves' disease;
KW   Hashimoto's thyroiditis; Wegener's granulomatosis; Omen's syndrome;
KW   chronic renal failure; autoimmune thyroid disease;
KW   acute infectious mononucleosis; HIV; herpes virus associated disease;
KW   common cold; human rhinovirus; coronavirus; enterovirus; herpes virus;
KW   influenza virus; parainfluenza virus; respiratory syncytial virus;
KW   adenovirus infection; bacteria pneumonia; sepsis; cerebral stroke;
KW   cerebral edema; ischaemia-reperfusion injury; hepatitis C; thrombolysis;
KW   cardiopulmonary bypass; percutaneous coronary intervention; PCT;
KW   coronary artery bypass; cardiac transplantation;
KW   isolated cerebral angiitis; mononeuritis multiplex;
KW   acute myocardial infarction; myocarditis; pericarditis;
KW   Liebman-Sachs endocarditis; chronic obstructive pulmonary disease; COPD;
KW   alveolitis; obliterating bronchiolitis; cystic fibrosis;
KW   allergic aspergillosis; Lofflers syndrome; sclerosing cholangiolitis;
KW   chronic cystitis; tubulo-interstitial nephritis;
KW   severe acute respiratory syndrome; SARS; large vessel vasculitis;
KW   giant cell arteritis; polymyalgia rheumatica; Takayasu arteritis;
KW   medium-sized vessel vasculitis; polyarteritis nodosa;
KW   localized polyarteritis nodosa; Kawasaki disease;
KW   small vessel vasculitis; Churg-Strauss syndrome;
KW   microscopic polyarteritis; cryoglobulinaemic vasculitis;
KW   leucocytoclastic angiitis; secondary vasculitis; rheumatoid vasculitis;
KW   systemic lupus erythematosus; isolated sacroileitis; SAPHO syndrome;
KW   disciitis; postoperative disciitis; subacute thyroiditis;
KW   cicatricial pemphigoid; dermatitis herpetiformis;
KW   subcorneal pustular dermatosis; epidermolysis bullosa acquisita; rosacea;
KW   acute febrile dermatosis; granuloma amiulare; Sweet's syndrome;
KW   pyoderma gangraenosum; acne; acne conglobata; sarcoidosis;
KW   relapsing polychondritis; familial Mediterranean fever; panniculitis;
KW   erythema nodosum; Weber-Christian's disease; retroperitoneal fibrosis;
KW   osteoporosis; osteolytic metastases; gastric cancer; colorectal cancer;
KW   urine bladder cancer; tumour growth; melanoma; thyroid carcinoma;
KW   transitional cell carcinoma; trichilemmona; squamous cell carcinoma;
KW   breast cancer; ss; gene.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   CDS             1. .342
FT                   /*tag=  a
FT                   /product= "VH 3-33 "
XX
PN   WO2004058797-A2.
XX
PD   15-JUL-2004.
XX
PF   16-DEC-2003; 2003WO-US040039.
XX
PR   16-DEC-2002; 2002US-0433728P.
XX
PA   (MEDA-) MEDAREX INC.
PA   (GENM-) GENMAB AS.
XX
PI   Teeling J,  Parren P,  Baadsgaard ODMS,  Hudson D,  Petersen J;
XX
DR   WPI; 2004-534122/51.
DR   P-PSDB; ADQ75776.
XX
PT   New isolated human monoclonal antibodies binding to human IL-8 having a
PT   heavy and/or light chain encoded by a nucleic acid, useful for preventing
PT   or treating cancer, immune disorders, inflammatory diseases and
PT   angiogenic conditions.
XX
PS   Claim 22; SEQ ID NO 9; 102pp; English.
XX
```

CC This sequence encodes the heavy chain variable region (VH) of the
CC isolated human monoclonal antibody (MAb), VH 3-33, which may be used as a
CC framework in the generation of the MAb of the invention which binds to
CC human interleukin (IL)-8. The MAb of the invention are useful for the
CC prevention and/or treatment of an inflammatory or hyperproliferative skin
CC disorder selected from PPP, psoriasis, including plaque psoriasis and
CC guttate type psoriasis, bullous skin diseases, such as bullous
CC pemphigoid, contact dermatitis, eczema, erythematosus, and atopic
CC dermatitis. The disorder is also an immune, autoimmune or inflammatory
CC disease selected from the group consisting of psoriatic arthritis,
CC systemic scleroderma and sclerosis, inflammatory bowel disease (IBD),
CC Crohn's disease, ulcerative colitis, acute lung injury, such as acute
CC respiratory distress syndrome or adult respiratory distress syndrome,
CC meningitis, encephalitis, uveitis, multiple myeloma, glomerulonephritis,
CC nephritis, asthma, atherosclerosis, leukocyte adhesion deficiency,
CC multiple sclerosis, Raynaud's syndrome, Sjogren's syndrome, juvenile
CC onset diabetes, Reiter's disease, Behcet's disease, immune complex
CC nephritis, IgA nephropathy, IgM polyneuropathies, immune-mediated
CC thrombocytopenias, such as acute idiopathic thrombocytopenic purpura and
CC chronic idiopathic thrombocytopenic purpura, haemolytic anaemia,
CC myasthenia gravis, lupus nephritis, lupus erythematosus, rheumatoid
CC arthritis (RA), ankylosing spondylitis, pemphigus, Graves' disease,
CC Hashimoto's thyroiditis, small vessel vasculitides, such as Wegener's
CC granulomatosis, Omen's syndrome, chronic renal failure, autoimmune
CC thyroid disease, acute infectious mononucleosis, HIV, herpes virus
CC associated diseases, human virus infections, such as common cold as
CC caused by human rhinovirus, coronavirus, other enterovirus, herpes virus,
CC influenza virus, parainfluenza virus, respiratory syncytial virus or
CC adenovirus infection, bacteria pneumonia, wounds, sepsis, cerebral
CC stroke/cerebral edema, ischaemia-reperfusion injury, and hepatitis C,
CC where the ischaemia-reperfusion injury is after thrombolysis,
CC cardiopulmonary bypass, percutaneous coronary intervention (PCT),
CC coronary artery bypass, or cardiac transplantation. The disorder is also
CC selected from isolated cerebral angiitis, mononeuritis multiplex, acute
CC myocardial infarction, myocarditis, pericarditis, and Liebman-Sachs
CC endocarditis, chronic obstructive pulmonary disease (COPD), alveolitis,
CC obliterating bronchiolitis, cystic fibrosis, allergic aspergillosis,
CC Lofflers syndrome, sclerosing cholangiolitis, chronic cystitis, tubulo-
CC interstitial nephritis, severe acute respiratory syndrome (SARS), large
CC vessel vasculitides (including giant cell arteritis, polymyalgia
CC rheumatica, and Takayasu arteritis), medium-sized vessel vasculitides
CC (including polyarteritis nodosa, localized polyarteritis nodosa, and
CC Kawasaki disease), small vessel vasculitides (including Churg-Strauss
CC syndrome, microscopic polyarteritis, cryoglobulinaemic vasculitis, and
CC leucocytoclastic angiitis), secondary vasculitides (including rheumatoid
CC vasculitis, and vasculitis associated with systemic lupus erythematosus
CC or Sjogren's syndrome), isolated sacroileitis, the SAPHO syndrome,
CC disciitis (including postoperative disciitis), subacute thyroiditis,
CC cicatricial pemphigoid, dermatitis herpetiformis, subcorneal pustular
CC dermatosis, epidermolysis bullosa acquisita, rosacea, acute febrile
CC dermatosis, granuloma amiulare (including Sweet's syndrome), pyoderma
CC gangraenosum, acne (including acne conglobata), sarcoidosis, relapsing
CC polychondritis, familial Mediterranean fever, panniculitis, erythema
CC nodosum, Weber-Christian's disease, and retroperitoneal fibrosis,
CC osteoporosis, osteolytic metastases, gastric cancer, colorectal cancer,
CC and urine bladder cancer. They are also for inhibiting or preventing
CC tumour growth, treating or preventing cancer, such as melanoma, thyroid
CC carcinoma, transitional cell carcinoma, trichilemmona, squamous cell
CC carcinoma and breast cancer. The anti-idiotypic antibody is useful for
CC detecting the level of human monoclonal antibody against IL-8 in a
CC sample.
XX
SQ Sequence 342 BP; 77 A; 83 C; 108 G; 74 T; 0 U; 0 Other;

Query Match 85.3%; Score 289.2; DB 12; Length 342;
Best Local Similarity 92.6%; Pred. No. 4e-76;
Matches 315; Conservative 0; Mismatches 23; Indels 2; Gaps 1;

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Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | ||||||||||||||| |||||| |||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| |||||||||||||   |   |||| |||||||||||||||||||||||
Db         62 CCTGTGCAGCGTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||  |||||||||||||||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATGGTATGATGGAAGTAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGC--GAAAAAGGA 299
              |||||||||||||||||||| |||||||||||||||||||||||||  || | |
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGATACTT 301

Qy        300 AGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
              | |||||||||||||||||||||||||||||||||||||
Db        302 TGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 341
```

RESULT 5
ADI26655
ID ADI26655 standard; cDNA; 357 BP.
XX
AC ADI26655;
XX
DT 15-APR-2004 (first entry)
XX
DE Human anti IgM antibody sHIgM22 VH cDNA.
XX
KW Human; ss; antibody; IgM; remyelination; neuronal growth; autoantibody;
KW demyelination disease; multiple sclerosis; central nervous system; CNS;
KW axon; glial cell proliferation;
KW Theiler's murine encephalomyelitis virus infection; CNS injury;
KW spinal cord injury.
XX
OS Homo sapiens.
XX
PN US2003185827-A1.
XX
PD 02-OCT-2003.
XX
PF 13-NOV-2001; 2001US-00010729.
XX
PR 29-APR-1994; 94US-00236520.
PR 08-AUG-1996; 96US-00692084.
PR 07-JAN-1997; 97US-00779784.
PR 28-MAY-1999; 99US-00322862.
PR 30-MAY-2000; 2000US-00580787.
PR 05-DEC-2000; 2000US-00730473.
XX
PA (MAYO-) MAYO FOUND.
XX
PI Rodriguez M, Miller DJ, Pease LR;
XX
DR WPI; 2004-119219/12.
DR P-PSDB; ADI26654.
XX
PT New human immunoglobulin M antibody for treating or preventing a
PT demyelinating disease of the central nervous system in a human or
PT domestic animal, such as multiple sclerosis.
XX
PS Example 8; Fig 35; 159pp; English.
XX
CC The invention relates to an antibody (I) produced by injecting an
CC immunocompetent host with an antibody peptide, and harvesting the
CC antibody, where the peptide comprises a human anti-IgM antibody fragment
CC given in the specification, or active fragments. Also included are
CC stimulating remyelination of central nervous system (CNS) axons in a
CC mammal (comprising administering a monoclonal antibody, or mixtures,
CC monomers, active fragments, or recombinant antibodies derived from it,
CC characterised by their ability to bind structures and cells within the
CC CNS, including oligodendrocytes), stimulating the proliferation of glial

```
CC   cells in CNS axons in a mammal (comprising administering a monoclonal
CC   antibody, or mixtures, monomers, active fragments, or recombinant
CC   antibodies derived from it, characterised by their ability to bind
CC   structures and cells within the CNS), treating or preventing a
CC   demyelinating disease of the CNS in a mammal (comprising administering a
CC   monoclonal antibody, or mixtures, monomers, active fragments; or
CC   recombinant antibodies derived from it, characterised by their ability to
CC   bind structures and cells within the CNS, and to stimulate remyelination
CC   of axons of the CNS), stimulating, in vitro, the proliferation of glial
CC   cells from mixed cell culture, stimulating remyelination of CNS axons in
CC   a mammal, a DNA sequence (or degenerate variant of it) which encodes an
CC   antibody(or a peptide analogue, hapten, or active fragment of it, where
CC   the DNA sequence consists of a sequence encoding an anti IgM antibody), a
CC   probe capable of screening for the antibody, an assay for screening drugs
CC   and other agents for the ability to modulate the production or mimic the
CC   activities of mAB sHIgM22, sHIgM46, or combinations of them, a
CC   recombinant virus transformed with recombinant antibody nucleic acids or
CC   vector, imaging a portion of the CNS using the antibody and diagnosing or
CC   monitoring demyelination and/or remyelination of the CNS comprising using
CC   CNS image. The antibody is used to stimulate remyelination of CNS axons,
CC   and to stimulate the proliferation of glial cells in CNS axons,
CC   optionally in vitro. The antibody is used to treat or prevent a
CC   demyelinating disease of the CNS in a human or domestic animal, such as
CC   multiple sclerosis, or a disease, other injury or dysfunction of the CNS,
CC   preferably the mammal is a mouse infected with Strain DA of Theiler's
CC   murine encephalomyelitis virus. The antibody is used to treat a spinal
CC   cord injury and used to screen drugs and other agents for the ability to
CC   modulate the production or mimic the activities of the antibody. The
CC   antibody can be used to image a portion of the CNS which can be used to
CC   diagnose or monitor demyelination and/or remyelination of the CNS. The
CC   present sequence is a cDNA for a variable region of a human anti-IgM
CC   antibody.
XX
SQ   Sequence 357 BP; 81 A; 89 C; 107 G; 80 T; 0 U; 0 Other;

  Query Match             83.9%;  Score 284.4;  DB 12;  Length 357;
  Best Local Similarity   90.4%;  Pred. No. 1.1e-74;
  Matches  322;  Conservative    0;  Mismatches   16;  Indels   18;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||||| |||||| |||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||||||||||||||||||||||||||||||||||||||||||| |||
Db        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGAGGTGA 301

Qy        299 ----------------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                             | ||||||||||||||||||||||||||||||||||||||
Db        302 CTGCTATTCCCTACTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 357


RESULT 6
ADI22042
ID   ADI22042 standard; DNA; 351 BP.
XX
AC   ADI22042;
XX
DT   22-APR-2004  (first entry)
XX
DE   Anti-platelet autoantibody related heavy chain nucleotide H37 SEQ:5.
XX
KW   anti-platelet autoantibody; autoantibody; blood clotting inhibition;
KW   thrombus; platelet adhesion inhibition;
KW   thrombotic thrombocytopenic purpura; platelet aggregation inhibition;
KW   idiopathic thrombocytopenic purpura; haemostatic; anticoagulant;
KW   thrombolytic; human; gene; ds.
XX
OS   Homo sapiens.
OS   Synthetic.
XX
PN   WO2004005890-A2.
XX
PD   15-JAN-2004.
XX
PF   03-JUL-2003; 2003WO-US021304.
XX
PR   03-JUL-2002; 2002US-0394352P.
PR   18-SEP-2002; 2002US-0411694P.
XX
PA   (UYPE-) UNIV PENNSYLVANIA.
XX
PI   Siegel DL;
XX
DR   WPI; 2004-142998/14.
DR   P-PSDB; ADI22095.
XX
PS   Claim 21; SEQ ID NO 5; 232pp; English.
XX
CC   The present invention describes a method (M1) for identifying an anti-
CC   platelet autoantibody (I) in a mammal. The autoantibody is detected by
CC   producing an antibody phage display library from B-lymphocytes obtained
CC   from the mammal, and screening the library to detect a phage that
CC   specifically binds with a platelet component, where the screening
CC   comprises panning the phage on intact platelets using competitive cell-
CC   surface panning. Also described: (1) an autoantibody identified by (M1);
CC   (2) an isolated nucleic acid encoding an anti-platelet autoantibody; (3)
CC   inhibiting (M2) blood clotting in a mammal having a thrombus or at risk
CC   of thrombus formation; (4) reversibly (M3) inhibiting blood clotting in a
CC   mammal having a thrombus or at risk of thrombus formation; (5) inhibiting
CC   (M4) binding of an anti-platelet autoantibody with a platelet component;
CC   (6) inhibiting (M5) platelet adhesion in a mammal; (7) treating (M6)
CC   thrombotic thrombocytopenic purpura in a mammal; (8) inhibiting (M7)
CC   platelet aggregation; (9) inhibiting (M8) platelet activation; (10)
CC   inhibiting (M9) platelet function; (11) inhibiting (M10) binding of an
CC   anti-platelet autoantibody, or its biologically active fragment with a
CC   platelet; (12) identifying (M11) a peptide that inhibits binding of an
CC   anti-platelet autoantibody with a platelet; (13) a peptide identified by
CC   the method of (12); (14) a peptide that specifically binds with an anti-
CC   platelet autoantibody; (15) treating (M12) idiopathic thrombocytopenic
CC   purpura (ITP) in a mammal; and (16) a kit for reversibly inhibiting blood
CC   clotting, inhibiting platelet aggregation, inhibiting platelet function
CC   or inhibiting platelet activation comprising an amount of an anti-
CC   platelet autoantibody, or its biologically active fragment that
CC   specifically binds with glycoprotein IIb/IIIa, where the autoantibody, or
CC   its fragment comprises an antigen binding region derived from an H44L4
CC   anti-platelet autoantibody, the kit further comprising a peptide
CC   inhibitor of the binding with glycoprotein IIb/IIIa, and an applicator
CC   and an instructions for use. (I) has haemostatic, anticoagulant and
CC   thrombolytic activities. The autoantibodies (I) are useful for diagnosing
CC   and for developing therapeutics for diseases mediated by autoantibody
CC   binding with platelet antigens. (M6) and (M12) are useful for treating
CC   thrombotic thrombocytopenic purpura and idiopathic thrombocytopenic
CC   purpura, respectively. (M2) and (M3) are useful for inhibiting blood
CC   clotting. The present sequence is used in the exemplification of the
CC   present invention.
XX
SQ   Sequence 351 BP; 79 A; 87 C; 108 G; 77 T; 0 U; 0 Other;

  Query Match             83.8%;  Score 284;  DB 12;  Length 351;
  Best Local Similarity   90.9%;  Pred. No. 1.5e-74;
  Matches  318;  Conservative    0;  Mismatches   20;  Indels   12;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
```

```
                ||||||||||| | ||||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
               |||||||||||||||||||||||    | | ||||| ||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAACTATGCTATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
               |||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTACG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
               |||||||||||||||||| |||||||||||||||||||||||||||||||| |   |
Db        242 TGCAAATGAACAGCCTGAGACCTGAGGACACGGCTGTGTATTACTGTGCGAGAGGTGGGG 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | |||||||||||||||||||||||||||||||||||||
Db        302 TAGCAGCTTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351
```

RESULT 7
ADI22043
ID ADI22043 standard; DNA; 351 BP.
XX
AC ADI22043;
XX
DT 22-APR-2004 (first entry)
XX
DE Anti-platelet autoantibody related heavy chain nucleotide H38 SEQ:6.
XX
KW anti-platelet autoantibody; autoantibody; blood clotting inhibition;
KW thrombus; platelet adhesion inhibition;
KW thrombotic thrombocytopenic purpura; platelet aggregation inhibition;
KW idiopathic thrombocytopenic purpura; haemostatic; anticoagulant;
KW thrombolytic; human; gene; ds.
XX
OS Homo sapiens.
OS Synthetic.
XX
PN WO2004005890-A2.
XX
PD 15-JAN-2004.
XX
PF 03-JUL-2003; 2003WO-US021304.
XX
PR 03-JUL-2002; 2002US-0394352P.
PR 18-SEP-2002; 2002US-0411694P.
XX
PA (UYPE-) UNIV PENNSYLVANIA.
XX
PI Siegel DL;
XX
DR WPI; 2004-142998/14.
DR P-PSDB; ADI22096.
XX
PS Claim 21; SEQ ID NO 6; 232pp; English.
XX
CC The present invention describes a method (M1) for identifying an anti-
CC platelet autoantibody (I) in a mammal. The autoantibody is detected by
CC producing an antibody phage display library from B-lymphocytes obtained
CC from the mammal, and screening the library to detect a phage that
CC specifically binds with a platelet component, where the screening
CC comprises panning the phage on intact platelets using competitive cell-
CC surface panning. Also described: (1) an autoantibody identified by (M1);
CC (2) an isolated nucleic acid encoding an anti-platelet autoantibody; (3)
CC inhibiting (M2) blood clotting in a mammal having a thrombus or at risk
CC of thrombus formation; (4) reversibly (M3) inhibiting blood clotting in a
CC mammal having a thrombus or at risk of thrombus formation; (5) inhibiting
CC (M4) binding of an anti-platelet autoantibody with a platelet component;
CC (6) inhibiting (M5) platelet adhesion in a mammal; (7) treating (M6)
CC thrombotic thrombocytopenic purpura in a mammal; (8) inhibiting (M7)
CC platelet aggregation; (9) inhibiting (M8) platelet activation; (10)
CC inhibiting (M9) platelet function; (11) inhibiting (M10) binding of an
CC anti-platelet autoantibody, or its biologically active fragment with a
CC platelet; (12) identifying (M11) a peptide that inhibits binding of an
CC anti-platelet autoantibody with a platelet; (13) a peptide identified by
CC the method of (12); (14) a peptide that specifically binds with an anti-
CC platelet autoantibody; (15) treating (M12) idiopathic thrombocytopenic
CC purpura (ITP) in a mammal; and (16) a kit for reversibly inhibiting blood
CC clotting, inhibiting platelet aggregation, inhibiting platelet function
CC or inhibiting platelet activation comprising an amount of an anti-
CC platelet autoantibody, or its biologically active fragment that
CC specifically binds with glycoprotein IIb/IIIa, where the autoantibody, or
CC its fragment comprises an antigen binding region derived from an H44L4
CC anti-platelet autoantibody, the kit further comprising a peptide
CC inhibitor of the binding with glycoprotein IIb/IIIa, and an applicator
CC and an instructions for use. (I) has haemostatic, anticoagulant and
CC thrombolytic activities. The autoantibodies (I) are useful for diagnosing
CC and for developing therapeutics for diseases mediated by autoantibody
CC binding with platelet antigens. (M6) and (M12) are useful for treating
CC thrombotic thrombocytopenic purpura and idiopathic thrombocytopenic
CC purpura, respectively. (M2) and (M3) are useful for inhibiting blood
CC clotting. The present sequence is used in the exemplification of the
CC present invention.
XX
SQ Sequence 351 BP; 79 A; 86 C; 109 G; 77 T; 0 U; 0 Other;

  Query Match   83.8%; Score 284; DB 12; Length 351;
  Best Local Similarity   90.9%; Pred. No. 1.5e-74;
  Matches 318; Conservative   0; Mismatches   20; Indels   12; Gaps   1;

```
Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||    | | ||||| ||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAACTATGCTATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTACG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||||||||| |||||||||||||||||||||||||||||||| |   |
Db        242 TGCAAATGAACAGCCTGAGACCTGAGGACACGGCTGTGTATTACTGTGCGAGAGGTGGGG 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | |||||||||||||||||||||||||||||||||||||
Db        302 TAGGAGCTTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351
```

RESULT 8
ADC61031
ID ADC61031 standard; DNA; 405 BP.
XX
AC ADC61031;
XX
DT 18-DEC-2003 (first entry)
XX
DE Human anti-CD45RB monoclonal antibody DNA, SEQ ID No 58.
XX
KW monoclonal antibody; CD45RB; antigen; cell proliferation;
KW immunosuppressive; neuroprotective; tissue rejection; organ rejection;
KW autoimmune disease; multiple sclerosis; human; anti-CD45RB; gene; ds.
XX
OS Homo sapiens.
XX

PN WO2003048327-A2.
XX
PD 12-JUN-2003.
XX
PF 02-DEC-2002; 2002WO-US038540.
XX
PR 03-DEC-2001; 2001US-0337276P.
XX
PA (ABGE-) ABGENIX INC.
XX
PI Foltz I, Babcook J, Palathumpat V, Yang X, King CT;
XX
DR WPI; 2003-558954/52.
DR P-PSDB; ADC61032.
XX
PT New anti-CD45RB monoclonal antibody, useful for treating an autoimmune
PT disease e.g. multiple sclerosis.
XX
PS Disclosure; SEQ ID NO 58; 121pp; English.
XX
CC The invention relates to a novel isolated monoclonal antibody comprising
CC a heavy chain having a sequence chosen from one of 22 fully defined
CC sequences comprising 135-147 amino acids, given in the specification, and
CC is specific for CD45RB antigen. The invention further relates to:
CC inhibiting cell proliferation associated with the expression of CD45RB
CC antigen; and treating a disease associated with the expression of a
CC CD45RB antigen in a patient. The monoclonal antibody has the activities
CC of immunosuppressive and neuroprotective. The monoclonal antibody is
CC useful for treating the rejection of a mammalian cell, tissue or organ,
CC especially an autoimmune disease in a mammal, especially a human e.g.
CC multiple sclerosis. This polynucleotide sequence represents a DNA
CC encoding a human protein of the anti-CD45RB monoclonal antibody of the
CC invention.
XX
SQ Sequence 405 BP; 84 A; 91 C; 134 G; 96 T; 0 U; 0 Other;

Query Match 83.7%; Score 283.8; DB 10; Length 405;
Best Local Similarity 91.1%; Pred. No. 1.8e-74;
Matches 316; Conservative 0; Mismatches 22; Indels 9; Gaps 1;

```
Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||| || | |||||||||||||| |||||| |||||||||| ||||||||||||||
Db         59 AGGTGCAGTTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 118

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| ||||||||||||||    | | |||| ||||||||||||||||||||||||
Db        119 CCTGTGCAGCGTCTGGATTCACCTTCAGTAACTATGGCATGCACTGGGTCCGCCAGGCTC 178

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||  |||||||||||||||||||||||||
Db        179 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATGGTATGATGGAAGTAATAAATACTATG 238

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              |||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db        239 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCGAAGAACACGCTGTATC 298

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              |||||||||||||||||||| |||||||||||||||||||||||||| |  ||  |
Db        299 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGGGGGGG 358

Qy        302 G---------CTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
              |         ||||||||||||||||||||||||||||||||||||
Db        359 GTGACTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 405
```

RESULT 9
ADI22044
ID ADI22044 standard; DNA; 351 BP.
XX
AC ADI22044;
XX
DT 22-APR-2004 (first entry)
XX
DE Anti-platelet autoantibody related heavy chain nucleotide H39 SEQ:7.
XX
KW anti-platelet autoantibody; autoantibody; blood clotting inhibition;
KW thrombus; platelet adhesion inhibition;
KW thrombotic thrombocytopenic purpura; platelet aggregation inhibition;
KW idiopathic thrombocytopenic purpura; haemostatic; anticoagulant;
KW thrombolytic; human; gene; ds.
XX
OS Homo sapiens.
OS Synthetic.
XX
PN WO2004005890-A2.
XX
PD 15-JAN-2004.
XX
PF 03-JUL-2003; 2003WO-US021304.
XX
PR 03-JUL-2002; 2002US-0394352P.
PR 18-SEP-2002; 2002US-0411694P.
XX
PA (UYPE-) UNIV PENNSYLVANIA.
XX
PI Siegel DL;
XX
DR WPI; 2004-142998/14.
DR P-PSDB; ADI22097.
XX
PS Claim 21; SEQ ID NO 7; 232pp; English.
XX
CC The present invention describes a method (M1) for identifying an anti-
CC platelet autoantibody (I) in a mammal. The autoantibody is detected by
CC producing an antibody phage display library from B-lymphocytes obtained
CC from the mammal, and screening the library to detect a phage that
CC specifically binds with a platelet component, where the screening
CC comprises panning the phage on intact platelets using competitive cell-
CC surface panning. Also described: (1) an autoantibody identified by (M1);
CC (2) an isolated nucleic acid encoding an anti-platelet autoantibody; (3)
CC inhibiting (M2) blood clotting in a mammal having a thrombus or at risk
CC of thrombus formation; (4) reversibly (M3) inhibiting blood clotting in a
CC mammal having a thrombus or at risk of thrombus formation; (5) inhibiting
CC (M4) binding of an anti-platelet autoantibody with a platelet component;
CC (6) inhibiting (M5) platelet adhesion in a mammal; (7) treating (M6)
CC thrombotic thrombocytopenic purpura in a mammal; (8) inhibiting (M7)
CC platelet aggregation; (9) inhibiting (M8) platelet activation; (10)
CC inhibiting (M9) platelet function; (11) inhibiting (M10) binding of an
CC anti-platelet autoantibody, or its biologically active fragment with a
CC platelet; (12) identifying (M11) a peptide that inhibits binding of an
CC anti-platelet autoantibody with a platelet; (13) a peptide identified by
CC the method of (12); (14) a peptide that specifically binds with an anti-
CC platelet autoantibody; (15) treating (M12) idiopathic thrombocytopenic
CC purpura (ITP) in a mammal; and (16) a kit for reversibly inhibiting blood
CC clotting, inhibiting platelet aggregation, inhibiting platelet function
CC or inhibiting platelet activation comprising an amount of an anti-
CC platelet autoantibody, or its biologically active fragment that
CC specifically binds with glycoprotein IIb/IIIa, where the autoantibody, or
CC its fragment comprises an antigen binding region derived from an H44L4
CC anti-platelet autoantibody, the kit further comprising a peptide
CC inhibitor of the binding with glycoprotein IIb/IIIa, and an applicator
CC and an instructions for use. (I) has haemostatic, anticoagulant and
CC thrombolytic activities. The autoantibodies (I) are useful for diagnosing
CC and for developing therapeutics for diseases mediated by autoantibody
CC binding with platelet antigens. (M6) and (M12) are useful for treating
CC thrombotic thrombocytopenic purpura and idiopathic thrombocytopenic
CC purpura, respectively. (M2) and (M3) are useful for inhibiting blood
CC clotting. The present sequence is used in the exemplification of the
CC present invention.
XX
SQ Sequence 351 BP; 79 A; 86 C; 108 G; 78 T; 0 U; 0 Other;

Query Match 83.3%; Score 282.4; DB 12; Length 351;
Best Local Similarity 90.6%; Pred. No. 4.4e-74;
Matches 317; Conservative 0; Mismatches 21; Indels 12; Gaps 1;

```
Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    | | ||||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAACTATGCTATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTACG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||| |||||| |||||||||||||||||||||||||||||||| |   |
Db        242 TGCAAATGAACAGTCTGAGACCTGAGGACACGGCTGTGTATTACTGTGCGAGAGGTGGGG 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | |||||||||||||||||||||||||||||||||||||
Db        302 TAGCAGCTTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351
```

RESULT 10
ADI22041
ID ADI22041 standard; DNA; 351 BP.
XX
AC ADI22041;
XX
DT 22-APR-2004 (first entry)
XX
DE Anti-platelet autoantibody related heavy chain nucleotide H36 SEQ:4.
XX
KW anti-platelet autoantibody; autoantibody; blood clotting inhibition;
KW thrombus; platelet adhesion inhibition;
KW thrombotic thrombocytopenic purpura; platelet aggregation inhibition;
KW idiopathic thrombocytopenic purpura; haemostatic; anticoagulant;
KW thrombolytic; human; gene; ds.
XX
OS Homo sapiens.
OS Synthetic.
XX
PN WO2004005890-A2.
XX
PD 15-JAN-2004.
XX
PF 03-JUL-2003; 2003WO-US021304.
XX
PR 03-JUL-2002; 2002US-0394352P.
PR 18-SEP-2002; 2002US-0411694P.
XX
PA (UYPE-) UNIV PENNSYLVANIA.
XX
PI Siegel DL;
XX
DR WPI; 2004-142998/14.
DR P-PSDB; ADI22094.
XX
PS Claim 21; SEQ ID NO 4; 232pp; English.
XX
CC The present invention describes a method (M1) for identifying an anti-
CC platelet autoantibody (I) in a mammal. The autoantibody is detected by
CC producing an antibody phage display library from B-lymphocytes obtained
CC from the mammal, and screening the library to detect a phage that
CC specifically binds with a platelet component, where the screening
CC comprises panning the phage on intact platelets using competitive cell-
CC surface panning. Also described: (1) an autoantibody identified by (M1);
CC (2) an isolated nucleic acid encoding an anti-platelet autoantibody; (3)
CC inhibiting (M2) blood clotting in a mammal having a thrombus or at risk
CC of thrombus formation; (4) reversibly (M3) inhibiting blood clotting in a
CC mammal having a thrombus or at risk of thrombus formation; (5) inhibiting
CC (M4) binding of an anti-platelet autoantibody with a platelet component;
CC (6) inhibiting (M5) platelet adhesion in a mammal; (7) treating (M6)
CC thrombotic thrombocytopenic purpura in a mammal; (8) inhibiting (M7)
CC platelet aggregation; (9) inhibiting (M8) platelet activation; (10)
CC inhibiting (M9) platelet function; (11) inhibiting (M10) binding of an
CC anti-platelet autoantibody, or its biologically active fragment with a
CC platelet; (12) identifying (M11) a peptide that inhibits binding of an
CC anti-platelet autoantibody with a platelet; (13) a peptide identified by
CC the method of (12); (14) a peptide that specifically binds with an anti-
CC platelet autoantibody; (15) treating (M12) idiopathic thrombocytopenic
CC purpura (ITP) in a mammal; and (16) a kit for reversibly inhibiting blood
CC clotting, inhibiting platelet aggregation, inhibiting platelet function
CC or inhibiting platelet activation comprising an amount of an anti-
CC platelet autoantibody, or its biologically active fragment that
CC specifically binds with glycoprotein IIb/IIIa, where the autoantibody, or
CC its fragment comprises an antigen binding region derived from an H44L4
CC anti-platelet autoantibody, the kit further comprising a peptide
CC inhibitor of the binding with glycoprotein IIb/IIIa, and an applicator
CC and an instructions for use. (I) has haemostatic, anticoagulant and
CC thrombolytic activities. The autoantibodies (I) are useful for diagnosing
CC and for developing therapeutics for diseases mediated by autoantibody
CC binding with platelet antigens. (M6) and (M12) are useful for treating
CC thrombotic thrombocytopenic purpura and idiopathic thrombocytopenic
CC purpura, respectively. (M2) and (M3) are useful for inhibiting blood
CC clotting. The present sequence is used in the exemplification of the
CC present invention.
XX
SQ Sequence 351 BP; 78 A; 87 C; 109 G; 77 T; 0 U; 0 Other;

```
  Query Match             83.3%;  Score 282.4;  DB 12;  Length 351;
  Best Local Similarity   90.6%;  Pred. No. 4.4e-74;
  Matches  317;  Conservative    0;  Mismatches   21;  Indels   12;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |   ||||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGCTATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTACG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||||||||| |||||||||||||||||||||||||||||| |   |
Db        242 TGCAAATGAACAGCCTGAGACCTGAGGACACGGCTGTGTATTACTGTGCGAGAGGTGGGG 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | |||||||||||||||||||||||||||||||||||||
Db        302 TAGCAGCTTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351
```

RESULT 11
ADI22052
ID ADI22052 standard; DNA; 345 BP.
XX
AC ADI22052;
XX
DT 22-APR-2004 (first entry)
XX
DE Anti-platelet autoantibody related heavy chain nucleotide H48 SEQ:15.
XX
KW anti-platelet autoantibody; autoantibody; blood clotting inhibition;
KW thrombus; platelet adhesion inhibition;
KW thrombotic thrombocytopenic purpura; platelet aggregation inhibition;
KW idiopathic thrombocytopenic purpura; haemostatic; anticoagulant;
KW thrombolytic; human; gene; ds.

```
XX
OS   Homo sapiens.
OS   Synthetic.
XX
PN   WO2004005890-A2.
XX
PD   15-JAN-2004.
XX
PF   03-JUL-2003; 2003WO-US021304.
XX
PR   03-JUL-2002; 2002US-0394352P.
PR   18-SEP-2002; 2002US-0411694P.
XX
PA   (UYPE-) UNIV PENNSYLVANIA.
XX
PI   Siegel DL;
XX
DR   WPI; 2004-142998/14.
DR   P-PSDB; ADI22105.
XX
PS   Claim 21; SEQ ID NO 15; 232pp; English.
XX
CC   The present invention describes a method (M1) for identifying an anti-
CC   platelet autoantibody (I) in a mammal. The autoantibody is detected by
CC   producing an antibody phage display library from B-lymphocytes obtained
CC   from the mammal, and screening the library to detect a phage that
CC   specifically binds with a platelet component, where the screening
CC   comprises panning the phage on intact platelets using competitive cell-
CC   surface panning. Also described: (1) an autoantibody identified by (M1);
CC   (2) an isolated nucleic acid encoding an anti-platelet autoantibody; (3)
CC   inhibiting (M2) blood clotting in a mammal having a thrombus or at risk
CC   of thrombus formation; (4) reversibly (M3) inhibiting blood clotting in a
CC   mammal having a thrombus or at risk of thrombus formation; (5) inhibiting
CC   (M4) binding of an anti-platelet autoantibody with a platelet component;
CC   (6) inhibiting (M5) platelet adhesion in a mammal; (7) treating (M6)
CC   thrombotic thrombocytopenic purpura in a mammal; (8) inhibiting (M7)
CC   platelet aggregation; (9) inhibiting (M8) platelet activation; (10)
CC   inhibiting (M9) platelet function; (11) inhibiting (M10) binding of an
CC   anti-platelet autoantibody, or its biologically active fragment with a
CC   platelet; (12) identifying (M11) a peptide that inhibits binding of an
CC   anti-platelet autoantibody with a platelet; (13) a peptide identified by
CC   the method of (12); (14) a peptide that specifically binds with an anti-
CC   platelet autoantibody; (15) treating (M12) idiopathic thrombocytopenic
CC   purpura (ITP) in a mammal; and (16) a kit for reversibly inhibiting blood
CC   clotting, inhibiting platelet aggregation, inhibiting platelet function
CC   or inhibiting platelet activation comprising an amount of an anti-
CC   platelet autoantibody, or its biologically active fragment that
CC   specifically binds with glycoprotein IIb/IIIa, where the autoantibody, or
CC   its fragment comprises an antigen binding region derived from an H44L4
CC   anti-platelet autoantibody, the kit further comprising a peptide
CC   inhibitor of the binding with glycoprotein IIb/IIIa, and an applicator
CC   and an instructions for use. (I) has haemostatic, anticoagulant and
CC   thrombolytic activities. The autoantibodies (I) are useful for diagnosing
CC   and for developing therapeutics for diseases mediated by autoantibody
CC   binding with platelet antigens. (M6) and (M12) are useful for treating
CC   thrombotic thrombocytopenic purpura and idiopathic thrombocytopenic
CC   purpura, respectively. (M2) and (M3) are useful for inhibiting blood
CC   clotting. The present sequence is used in the exemplification of the
CC   present invention.
XX
SQ   Sequence 345 BP; 81 A; 86 C; 99 G; 79 T; 0 U; 0 Other;

  Query Match             83.2%;  Score 282;  DB 12;  Length 345;
  Best Local Similarity   91.0%;  Pred. No. 5.7e-74;
  Matches  313;  Conservative    0;  Mismatches   25;  Indels    6;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              ||||||||||||||||||||||| |    | |||||  ||| ||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCATCAGTAATTATCACATACACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||| |||||||||||||||||||||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCATTTATATCATATGATGGAAGTAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              |||||||||||||||||||||||||||||||||||||||||||||||||||||| ||| ||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGTTGTTTC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA- 300
              |||||||||||||||||| | |||||||||||||||||||||||||||||| |   || |
Db        242 TGCAAATGAACAGCCTGACAACTGAGGACACGGCTGTGTATTACTGTGCGATAGTGGGAC 301

Qy        301 -----GGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                  | |||| |||||||||||||||||||||||||||| |||
Db        302 CCTTTGACTACCGGGGCCAGGGAACCCTGGTCACCGTCTCTTCA 345


RESULT 12
ABK88453
ID   ABK88453 standard; cDNA; 354 BP.
XX
AC   ABK88453;
XX
DT   07-OCT-2002  (first entry)
XX
DE   Human anti-CD40 monoclonal antibody 9F7 VH.1 region cDNA.
XX
KW   Human; VH.1; heavy chain variable region; CD40; ss; gene;
KW   autoimmune disease; systemic lupus erythematosus; psoriasis;
KW   multiple sclerosis; inflammatory bowel disease; Crohn's disease;
KW   rheumatoid arthritis; organ rejection; lymphoma; non-Hodgkin's lymphoma;
KW   monoclonal antibody; B cell; glycoprotein; proliferation.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   CDS             1. .354
FT                   /*tag=  a
FT                   /product= "9F7 heavy chain variable region"
FT                   /partial
FT                   /note= "No start or stop codon shown"
XX
PN   WO200228904-A2.
XX
PD   11-APR-2002.
XX
PF   02-OCT-2001; 2001WO-US030857.
XX
PR   02-OCT-2000; 2000US-0237556P.
XX
PA   (CHIR ) CHIRON CORP.
XX
PI   Chu K,  Wang C,  Yoshihara C,  Donnelly JJ;
XX
DR   WPI; 2002-405169/43.
DR   P-PSDB; ABG30463.
XX
PT   A human anti-CD40 monoclonal antibody or fragment useful for inhibiting
PT   proliferation, growth or differentiation of a normal human B cells and
PT   treating autoimmune disease such as rheumatoid arthritis or systemic
PT   lupus erythematosus.
XX
PS   Claim 11; Fig 11; 75pp; English.
XX
CC   The invention relates to a human monoclonal antibody or fragment capable
CC   of specifically binding to a human CD40 antigen (a glycoprotein expressed
CC   on the surface of human B cells), where the antibody or fragment is free
CC   of significant agonistic activity when it binds to the CD40 antigen, and
CC   the growth or differentiation is inhibited. The fragments comprise the
CC   complementarity determining region (CDR) of the light and heavy chains of
CC   the monoclonal antibodies secreted by a hybridoma consisting of 15B8,
CC   20C4, 12D9, 9F7 and 13E4. Also included are the nucleic acids encoding
```

CC the antibody (or fragments). The antibodies or fragments are used for
CC inhibiting proliferation, growth or differentiation of a normal human B
CC cells and to inhibit antibody production by B cells. They may also be
CC useful for treating autoimmune diseases, such as systemic lupus
CC erythematosus, psoriasis, multiple sclerosis, inflammatory bowel disease
CC (Crohn's disease), rheumatoid arthritis, and lymphoma (especially Non-
CC Hodgkin's Lymphoma). The present sequence encodes the antibody 9F7 heavy
CC chain variable region VH.1
XX
SQ Sequence 354 BP; 80 A; 91 C; 107 G; 76 T; 0 U; 0 Other;

Query Match 82.9%; Score 281; DB 6; Length 354;
Best Local Similarity 90.1%; Pred. No. 1.1e-73;
Matches 318; Conservative 0; Mismatches 20; Indels 15; Gaps 1;

```
Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              ||||||||||||  |||||||||||||| ||||||  |||||||||  ||||||||||||||||
Db          2 AGGTGCAGCTGCAGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTATTAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              | ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CTGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              ||||||||||||||||||||||||||||||||||||||||| |||||||||| | |  | |
Db        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTACTACTGTGCGAGAGATCATG 301

Qy        302 G---------------CTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
              |               ||||||||||||||||||||||||||||||||||||
Db        302 GGAGTAACCCCCTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 354
```

RESULT 13
ADI22046
ID ADI22046 standard; DNA; 351 BP.
XX
AC ADI22046;
XX
DT 22-APR-2004 (first entry)
XX
DE Anti-platelet autoantibody related heavy chain nucleotide H41 SEQ:9.
XX
KW anti-platelet autoantibody; autoantibody; blood clotting inhibition;
KW thrombus; platelet adhesion inhibition;
KW thrombotic thrombocytopenic purpura; platelet aggregation inhibition;
KW idiopathic thrombocytopenic purpura; haemostatic; anticoagulant;
KW thrombolytic; human; gene; ds.
XX
OS Homo sapiens.
OS Synthetic.
XX
PN WO2004005890-A2.
XX
PD 15-JAN-2004.
XX
PF 03-JUL-2003; 2003WO-US021304.
XX
PR 03-JUL-2002; 2002US-0394352P.
PR 18-SEP-2002; 2002US-0411694P.
XX
PA (UYPE-) UNIV PENNSYLVANIA.
XX
PI Siegel DL;
XX
DR WPI; 2004-142998/14.
DR P-PSDB; ADI22099.
XX
PS Claim 21; SEQ ID NO 9; 232pp; English.
XX
CC The present invention describes a method (M1) for identifying an anti-
CC platelet autoantibody (I) in a mammal. The autoantibody is detected by
CC producing an antibody phage display library from B-lymphocytes obtained
CC from the mammal, and screening the library to detect a phage that
CC specifically binds with a platelet component, where the screening
CC comprises panning the phage on intact platelets using competitive cell-
CC surface panning. Also described: (1) an autoantibody identified by (M1);
CC (2) an isolated nucleic acid encoding an anti-platelet autoantibody; (3)
CC inhibiting (M2) blood clotting in a mammal having a thrombus or at risk
CC of thrombus formation; (4) reversibly (M3) inhibiting blood clotting in a
CC mammal having a thrombus or at risk of thrombus formation; (5) inhibiting
CC (M4) binding of an anti-platelet autoantibody with a platelet component;
CC (6) inhibiting (M5) platelet adhesion in a mammal; (7) treating (M6)
CC thrombotic thrombocytopenic purpura in a mammal; (8) inhibiting (M7)
CC platelet aggregation; (9) inhibiting (M8) platelet activation; (10)
CC inhibiting (M9) platelet function; (11) inhibiting (M10) binding of an
CC anti-platelet autoantibody, or its biologically active fragment with a
CC platelet; (12) identifying (M11) a peptide that inhibits binding of an
CC anti-platelet autoantibody with a platelet; (13) a peptide identified by
CC the method of (12); (14) a peptide that specifically binds with an anti-
CC platelet autoantibody; (15) treating (M12) idiopathic thrombocytopenic
CC purpura (ITP) in a mammal; and (16) a kit for reversibly inhibiting blood
CC clotting, inhibiting platelet aggregation, inhibiting platelet function
CC or inhibiting platelet activation comprising an amount of an anti-
CC platelet autoantibody, or its biologically active fragment that
CC specifically binds with glycoprotein IIb/IIIa, where the autoantibody, or
CC its fragment comprises an antigen binding region derived from an H44L4
CC anti-platelet autoantibody, the kit further comprising a peptide
CC inhibitor of the binding with glycoprotein IIb/IIIa, and an applicator
CC and an instructions for use. (I) has haemostatic, anticoagulant and
CC thrombolytic activities. The autoantibodies (I) are useful for diagnosing
CC and for developing therapeutics for diseases mediated by autoantibody
CC binding with platelet antigens. (M6) and (M12) are useful for treating
CC thrombotic thrombocytopenic purpura and idiopathic thrombocytopenic
CC purpura, respectively. (M2) and (M3) are useful for inhibiting blood
CC clotting. The present sequence is used in the exemplification of the
CC present invention.
XX
SQ Sequence 351 BP; 78 A; 87 C; 108 G; 78 T; 0 U; 0 Other;

Query Match 82.8%; Score 280.8; DB 12; Length 351;
Best Local Similarity 90.3%; Pred. No. 1.3e-73;
Matches 316; Conservative 0; Mismatches 22; Indels 12; Gaps 1;

```
Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||  | ||||||||||||| |||||| |||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||| ||||||    | | ||||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATCCACCTTCAGTTACTATGCTATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTACG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||||||||||| |||||||||||||||||||||||||||||||| |   |
Db        242 TGCAAATGAACAGCCTGAGACCTGAGGACACGGCTGTGTATTACTGTGCGAGAGGTGGGG 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | |||||||||  ||||||||||||||||||||||||||||
Db        302 TAGCAGCTTTTGACTACTGGGGTCAGGGAACCCTGGTCACCGTCTCCTCA 351
```

RESULT 14
ADI22045
ID ADI22045 standard; DNA; 351 BP.
XX
AC ADI22045;
XX
DT 22-APR-2004 (first entry)
XX
DE Anti-platelet autoantibody related heavy chain nucleotide H40 SEQ:8.
XX
KW anti-platelet autoantibody; autoantibody; blood clotting inhibition;
KW thrombus; platelet adhesion inhibition;
KW thrombotic thrombocytopenic purpura; platelet aggregation inhibition;
KW idiopathic thrombocytopenic purpura; haemostatic; anticoagulant;
KW thrombolytic; human; gene; ds.
XX
OS Homo sapiens.
OS Synthetic.
XX
PN WO2004005890-A2.
XX
PD 15-JAN-2004.
XX
PF 03-JUL-2003; 2003WO-US021304.
XX
PR 03-JUL-2002; 2002US-0394352P.
PR 18-SEP-2002; 2002US-0411694P.
XX
PA (UYPE-) UNIV PENNSYLVANIA.
XX
PI Siegel DL;
XX
DR WPI; 2004-142998/14.
DR P-PSDB; ADI22098.
XX
PS Claim 21; SEQ ID NO 8; 232pp; English.
XX
CC The present invention describes a method (M1) for identifying an anti-
CC platelet autoantibody (I) in a mammal. The autoantibody is detected by
CC producing an antibody phage display library from B-lymphocytes obtained
CC from the mammal, and screening the library to detect a phage that
CC specifically binds with a platelet component, where the screening
CC comprises panning the phage on intact platelets using competitive cell-
CC surface panning. Also described: (1) an autoantibody identified by (M1);
CC (2) an isolated nucleic acid encoding an anti-platelet autoantibody; (3)
CC inhibiting (M2) blood clotting in a mammal having a thrombus or at risk
CC of thrombus formation; (4) reversibly (M3) inhibiting blood clotting in a
CC mammal having a thrombus or at risk of thrombus formation; (5) inhibiting
CC (M4) binding of an anti-platelet autoantibody with a platelet component;
CC (6) inhibiting (M5) platelet adhesion in a mammal; (7) treating (M6)
CC thrombotic thrombocytopenic purpura in a mammal; (8) inhibiting (M7)
CC platelet aggregation; (9) inhibiting (M8) platelet activation; (10)
CC inhibiting (M9) platelet function; (11) inhibiting (M10) binding of an
CC anti-platelet autoantibody, or its biologically active fragment with a
CC platelet; (12) identifying (M11) a peptide that inhibits binding of an
CC anti-platelet autoantibody with a platelet; (13) a peptide identified by
CC the method of (12); (14) a peptide that specifically binds with an anti-
CC platelet autoantibody; (15) treating (M12) idiopathic thrombocytopenic
CC purpura (ITP) in a mammal; and (16) a kit for reversibly inhibiting blood
CC clotting, inhibiting platelet aggregation, inhibiting platelet function
CC or inhibiting platelet activation comprising an amount of an anti-
CC platelet autoantibody, or its biologically active fragment that
CC specifically binds with glycoprotein IIb/IIIa, where the autoantibody, or
CC its fragment comprises an antigen binding region derived from an H44L4
CC anti-platelet autoantibody, the kit further comprising a peptide
CC inhibitor of the binding with glycoprotein IIb/IIIa, and an applicator
CC and an instructions for use. (I) has haemostatic, anticoagulant and
CC thrombolytic activities. The autoantibodies (I) are useful for diagnosing
CC and for developing therapeutics for diseases mediated by autoantibody
CC binding with platelet antigens. (M6) and (M12) are useful for treating
CC thrombotic thrombocytopenic purpura and idiopathic thrombocytopenic
CC purpura, respectively. (M2) and (M3) are useful for inhibiting blood
CC clotting. The present sequence is used in the exemplification of the
CC present invention.
XX
SQ Sequence 351 BP; 79 A; 88 C; 107 G; 77 T; 0 U; 0 Other;

Query Match 82.8%; Score 280.8; DB 12; Length 351;
Best Local Similarity 90.3%; Pred. No. 1.3e-73;
Matches 316; Conservative 0; Mismatches 22; Indels 12; Gaps 1;

```
Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              ||||||||||||||  |||||||||    | | ||||| ||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTACATTCACCTTCAGTAACTATGCTATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTACG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||||||||| |||||||||||||||||||||||||||||||||| |   |
Db        242 TGCAAATGAACAGCCTGAGACCTGAGGACACGGCTGTGTATTACTGTGCGAGAGGTGGGG 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | ||||||||||||||||||||||||||||||||||||||
Db        302 TAGCGGCTTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351
```

RESULT 15
ADS84379
ID ADS84379 standard; DNA; 349 BP.
XX
AC ADS84379;
XX
DT 18-NOV-2004 (first entry)
XX
DE Human anti-EPO-R antibody heavy chain variable region DNA SEQ ID NO:18.
XX
KW human; erythropoietin receptor; EPO receptor;
KW erythropoietin receptor binding antibody; EPO receptor binding antibody;
KW antianaemic; neuroprotective; vulnerary; gene therapy; aplasia; anaemia;
KW wound healing; neural cell damage protection;
KW neural tissue damage protection; brain injury; spinal cord injury;
KW stroke; anti-erythropoietin receptor antibody; anti-EPO-R antibody; gene;
KW ds.
XX
OS Homo sapiens.
XX
PN WO2004035603-A2.
XX
PD 29-APR-2004.
XX
PF 14-OCT-2003; 2003WO-US032243.
XX
PR 14-OCT-2002; 2002US-00269711.
PR 10-OCT-2003; 2003US-00684109.
XX
PA (ABBO) ABBOTT LAB.
XX
PI Devries PJ, Green LL, Ostrow DH, Reilly EB, Wieler J;
XX
DR WPI; 2004-348433/32.
DR P-PSDB; ADS84380.
XX
PT New antibodies that bind to or activate an endogenous human
PT erythropoietin receptor, useful for diagnosing, preventing or treating
PT disorders associated with dysfunctional erythropoietin receptor, e.g.
PT anemia.

```
XX
PS   Claim 47; SEQ ID NO 18; 192pp; English.
XX
CC   The present invention describes an antibody or its fragment that binds to
CC   or activates an endogenous activity of a human erythropoietin (EPO)
CC   receptor in a mammal, but does not interact with a peptide having a
CC   sequence of 30 amino acids (SEQ ID NO:1, ADS84362). Also described: (1)
CC   methods of modulating or activating an endogenous activity of a human EPO
CC   receptor in a mammal, comprising administering to the mammal a
CC   therapeutic amount of the above antibody or its fragment to modulate or
CC   activate the receptor; (2) a method of treating a mammal suffering from
CC   aplasia, comprising administering to the mammal a therapeutic amount of
CC   the above antibody or its fragment to modulate or activate the receptor;
CC   (3) a pharmaceutical composition comprising a therapeutic amount of the
CC   above antibody or antibody fragment, and a pharmaceutical excipient; (4)
CC   an isolated and purified polynucleotide sequence, and their fragments,
CC   complements and degenerate codon equivalents; and (5) an isolated and
CC   purified amino acid sequence, and their fragments. The EPO receptor
CC   binding antibody has antianaemic, neuroprotective and vulnerary
CC   activities, and can be used in gene therapy. The compositions and methods
CC   from the present invention can be used for modulating an endogenous
CC   activity of a human EPO receptor or for treating mammals suffering from
CC   aplasia or anaemia. They may also be used for identifying mammals having
CC   a dysfunctional EPO receptor. The composition may also be used in
CC   promoting wound healing or in protecting against neural cell and/or
CC   tissue damage resulting from brain/spinal cord injury, stroke and the
CC   like. The present sequence encodes a human anti-EPO-R antibody heavy
CC   chain variable region, which is given in the exemplification of the
CC   present invention.
XX
SQ   Sequence 349 BP; 78 A; 85 C; 110 G; 76 T; 0 U; 0 Other;

  Query Match             82.8%;  Score 280.6;  DB 13;  Length 349;
  Best Local Similarity   90.5%;  Pred. No. 1.5e-73;
  Matches  314;  Conservative    0;  Mismatches   24;  Indels    9;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||  | |||||  |||||||  ||||||  |||||||||  ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCGGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              ||||||||||  ||||||||||||||    | |  ||||  ||||||||||||||||||||||||
Db         62 CCTGTGCAGCGTCTGGATTCACCTTCAGTAAATATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||  |||  ||||||||||||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTTTATGGTATGATGGAAGTAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGA--------- 292
              ||||||||||||||||||||  ||||||||||||||||||||||||||
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGGTCCGT 301

Qy        293 AAAAGGAAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
              |  |    | ||||||||||||||||||||||||||||||||||||
Db        302 ACTACTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 348


Search completed: September 11, 2005, 20:25:57
Job time : 360.464 secs
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: September 11, 2005, 18:08:48 ; Search time 2724.33 Seconds
(without alignments)
4736.504 Million cell updates/sec

Title: US-09-403-107-145_COPY_1_339
Perfect score: 339
Sequence: 1 gaggtgcagctgctcgagtc.....ccctggtcaccgtctcctca 339

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 34239544 seqs, 19032134700 residues

Total number of hits satisfying chosen parameters: 68479088

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : EST:*
1: gb_est1:*
2: gb_est2:*
3: gb_htc:*
4: gb_est3:*
5: gb_est4:*
6: gb_est5:*
7: gb_est6:*
8: gb_gss1:*
9: gb_gss2:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	274.2	80.9	678	4	BG340648	BG340648 602462226
2	274.2	80.9	788	4	BG342203	BG342203 602462979
3	274.2	80.9	870	4	BG759649	BG759649 602713342
4	274.2	80.9	964	2	BF663281	BF663281 602144406
5	274.2	80.9	991	4	BF974524	BF974524 602243430
6	274.2	80.9	1010	2	BF663436	BF663436 602144593
7	273.2	80.6	846	4	BG755572	BG755572 602716255
8	272.6	80.4	912	4	BG756211	BG756211 602713521
9	271.8	80.2	413	5	BX344075	BX344075 BX344075
10	270.6	79.8	516	2	AW401728	AW401728 UI-HF-BK0
11	269.6	79.5	899	5	BQ420418	BQ420418 AGENCOURT
12	269.4	79.5	663	4	BG686759	BG686759 602650729
13	267.2	78.8	503	2	AW402572	AW402572 UI-HF-BK0
14	266.8	78.7	894	5	BQ708235	BQ708235 AGENCOURT
15	266.6	78.6	456	4	BG503730	BG503730 602549705
16	266.6	78.6	902	5	BX436850	BX436850 BX436850
17	266.4	78.6	1228	5	BM920476	BM920476 AGENCOURT
18	266.2	78.5	761	4	BG757080	BG757080 602715111
19	265.8	78.4	413	5	BX399298	BX399298 BX399298
20	264.8	78.1	918	5	BQ710537	BQ710537 AGENCOURT
21	264.6	78.1	621	6	CD687626	CD687626 EST4147 h
22	263.6	77.8	436	2	AW402311	AW402311 UI-HF-BK0
23	263.2	77.6	750	4	BF976111	BF976111 602244955
24	262.4	77.4	430	5	BX480277	BX480277 DKFZp686O
25	261.8	77.2	950	5	BQ708952	BQ708952 AGENCOURT
26	261.4	77.1	536	2	AW402624	AW402624 UI-HF-BK0
27	260.8	76.9	907	5	BQ708724	BQ708724 AGENCOURT
28	260.8	76.9	1001	5	BM914366	BM914366 AGENCOURT
29	259.8	76.6	891	5	BU150876	BU150876 AGENCOURT
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31	259.2	76.5	930	4	BG755605	BG755605 602716292
32	258.6	76.3	802	4	BG756245	BG756245 602713557
33	257	75.8	999	5	BQ707443	BQ707443 AGENCOURT
34	255.2	75.3	415	5	BX456327	BX456327 BX456327
35	254.6	75.1	511	2	AW402453	AW402453 UI-HF-BK0
36	253.8	74.9	937	5	BQ706852	BQ706852 AGENCOURT
37	253.2	74.7	434	5	BX480153	BX480153 DKFZp686C
38	253	74.6	405	2	AW402449	AW402449 UI-HF-BK0
39	253	74.6	962	5	BQ706534	BQ706534 AGENCOURT
40	253	74.6	1000	5	BQ712690	BQ712690 AGENCOURT
41	251.6	74.2	891	5	BQ709554	BQ709554 AGENCOURT
42	251	74.0	889	5	BQ709105	BQ709105 AGENCOURT
43	251	74.0	925	5	BQ706858	BQ706858 AGENCOURT
44	251	74.0	986	5	BQ710656	BQ710656 AGENCOURT
45	250.4	73.9	648	4	BG340670	BG340670 602462250

ALIGNMENTS

RESULT 1
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LOCUS BG340648 678 bp mRNA linear EST 27-FEB-2001
DEFINITION 602462226F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4574935 5', mRNA sequence.
ACCESSION BG340648
VERSION BG340648.1 GI:13147086
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 678)
AUTHORS NIH-MGC http://mgc.nci.nih.gov/.
TITLE National Institutes of Health, Mammalian Gene Collection (MGC)
JOURNAL Unpublished (1999)
COMMENT Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1285 row: d column: 08
High quality sequence stop: 666.
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/organism="Homo sapiens"
/mol_type="mRNA"
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/clone="IMAGE:4574935"
/tissue_type="primary B-cells from tonsils (cell line)"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_48"
/note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."
ORIGIN

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  Query Match             80.9%;  Score 274.2;  DB 4;  Length 678;
  Best Local Similarity   95.6%;  Pred. No. 1.2e-71;
  Matches  282;  Conservative    0;  Mismatches   13;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Db        190 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 249

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        250 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 309

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Db        310 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 369

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA 296
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Db        370 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGA 424


RESULT 2
BG342203
LOCUS       BG342203                 788 bp    mRNA    linear   EST 27-FEB-2001
DEFINITION  602462979F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4575931 5',
            mRNA sequence.
ACCESSION   BG342203
VERSION     BG342203.1  GI:13148641
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 788)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1287  row: m  column: 20
            High quality sequence stop: 687.
FEATURES             Location/Qualifiers
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                     /db_xref="taxon:9606"
                     /clone="IMAGE:4575931"
                     /tissue_type="primary B-cells from tonsils (cell line)"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_48"
                     /note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
                     Site_2: EcoRI; cDNA made by oligo-dT priming.
                     Directionally cloned into EcoRI/XhoI sites using the
                     following 5' adaptor: GGCACGAG(G). Size-selected >500bp
                     for average insert size 1.8kb. Library constructed by Ling
                     Hong in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using  ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             80.9%;  Score 274.2;  DB 4;  Length 788;
  Best Local Similarity   95.6%;  Pred. No. 1.2e-71;
  Matches  282;  Conservative    0;  Mismatches   13;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||| |||||| |||||||| ||||||||||||||||
Db        129 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 188

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Db        189 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 248

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Db        309 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 368

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA 296
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Db        369 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGA 423


RESULT 3
BG759649
LOCUS       BG759649                 870 bp    mRNA    linear   EST 15-MAY-2001
DEFINITION  602713342F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4853338 5',
            mRNA sequence.
ACCESSION   BG759649
VERSION     BG759649.1  GI:14070302
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 870)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1699  row: d  column: 11
            High quality sequence stop: 764.
FEATURES             Location/Qualifiers
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                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:4853338"
                     /tissue_type="primary B-cells from tonsils (cell line)"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_48"
                     /note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
                     Site_2: EcoRI; cDNA made by oligo-dT priming.
                     Directionally cloned into EcoRI/XhoI sites using the
                     following 5' adaptor: GGCACGAG(G). Size-selected >500bp
                     for average insert size 1.8kb. Library constructed by Ling
                     Hong in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using  ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             80.9%;  Score 274.2;  DB 4;  Length 870;
  Best Local Similarity   95.6%;  Pred. No. 1.2e-71;
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  Matches  282;  Conservative    0;  Mismatches   13;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Db        130 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 189

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA 296
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Db        370 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGA 424


RESULT 4
BF663281
LOCUS       BF663281                964 bp    mRNA    linear   EST 21-DEC-2000
DEFINITION  602144406F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4297849 5',
            mRNA sequence.
ACCESSION   BF663281
VERSION     BF663281.1  GI:11937163
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 964)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1152  row: k  column: 02
            High quality sequence stop: 693.
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                     /clone="IMAGE:4297849"
                     /tissue_type="primary B-cells from tonsils (cell line)"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_48"
                     /note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
                     Site_2: EcoRI; cDNA made by oligo-dT priming.
                     Directionally cloned into EcoRI/XhoI sites using the
                     following 5' adaptor: GGCACGAG(G). Size-selected >500bp
                     for average insert size 1.8kb. Library constructed by Ling
                     Hong in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using  ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             80.9%;  Score 274.2;  DB 2;  Length 964;
  Best Local Similarity   95.6%;  Pred. No. 1.3e-71;
  Matches  282;  Conservative    0;  Mismatches   13;  Indels    0;  Gaps    0;

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Db        190 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 249

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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA 296
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Db        370 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGA 424


RESULT 5
BF974524
LOCUS       BF974524                991 bp    mRNA    linear   EST 22-JAN-2001
DEFINITION  602243430F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4334719 5',
            mRNA sequence.
ACCESSION   BF974524
VERSION     BF974524.1  GI:12341739
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 991)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1204  row: k  column: 08
            High quality sequence stop: 688.
FEATURES             Location/Qualifiers
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                     /mol_type="mRNA"
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                     /clone="IMAGE:4334719"
                     /tissue_type="primary B-cells from tonsils (cell line)"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_48"
                     /note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
                     Site_2: EcoRI; cDNA made by oligo-dT priming.
                     Directionally cloned into EcoRI/XhoI sites using the
                     following 5' adaptor: GGCACGAG(G). Size-selected >500bp
                     for average insert size 1.8kb. Library constructed by Ling
                     Hong in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using  ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             80.9%;  Score 274.2;  DB 4;  Length 991;
  Best Local Similarity   95.6%;  Pred. No. 1.3e-71;
  Matches  282;  Conservative    0;  Mismatches   13;  Indels    0;  Gaps    0;
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Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Db        130 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 189

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA 296
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Db        370 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGA 424


RESULT 6
BF663436
LOCUS       BF663436                1010 bp    mRNA    linear   EST 21-DEC-2000
DEFINITION  602144593F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4297847 5',
            mRNA sequence.
ACCESSION   BF663436
VERSION     BF663436.1  GI:11937331
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1010)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1152  row: j  column: 24
            High quality sequence stop: 695.
FEATURES             Location/Qualifiers
     source          1. .1010
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:4297847"
                     /tissue_type="primary B-cells from tonsils (cell line)"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_48"
                     /note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
                     Site_2: EcoRI; cDNA made by oligo-dT priming.
                     Directionally cloned into EcoRI/XhoI sites using the
                     following 5' adaptor: GGCACGAG(G). Size-selected >500bp
                     for average insert size 1.8kb. Library constructed by Ling
                     Hong in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using  ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             80.9%;  Score 274.2;  DB 2;  Length 1010;
  Best Local Similarity   95.6%;  Pred. No. 1.3e-71;
  Matches  282;  Conservative    0;  Mismatches   13;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||||
Db        130 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 189

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |   |||| ||||||||||||||||||||||||
Db        190 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 249

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        250 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 309

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        310 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 369

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA 296
              |||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        370 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGA 424


RESULT 7
BG755572
LOCUS       BG755572                846 bp    mRNA    linear   EST 15-MAY-2001
DEFINITION  602716255F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4856628 5',
            mRNA sequence.
ACCESSION   BG755572
VERSION     BG755572.1  GI:14066225
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 846)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1707  row: m  column: 13
            High quality sequence stop: 752.
FEATURES             Location/Qualifiers
     source          1. .846
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:4856628"
                     /tissue_type="primary B-cells from tonsils (cell line)"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_48"
                     /note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
                     Site_2: EcoRI; cDNA made by oligo-dT priming.
                     Directionally cloned into EcoRI/XhoI sites using the
                     following 5' adaptor: GGCACGAG(G). Size-selected >500bp
                     for average insert size 1.8kb. Library constructed by Ling
                     Hong in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using  ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             80.6%;  Score 273.2;  DB 4;  Length 846;
  Best Local Similarity   95.6%;  Pred. No. 2.5e-71;
  Matches  281;  Conservative    0;  Mismatches   13;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||||
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Db        130 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 189

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              ||||||||||||||||||||||||    |   |||| ||||||||||||||||||||||||
Db        190 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 249

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        250 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 309

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        310 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 369

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAA 295
              |||||||||||||||||||||||||||||||||||||||||||||||||||| ||
Db        370 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGACAA 423


RESULT 8
BG756211
LOCUS       BG756211                 912 bp    mRNA    linear   EST 15-MAY-2001
DEFINITION  602713521F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4853837 5',
            mRNA sequence.
ACCESSION   BG756211
VERSION     BG756211.1  GI:14066864
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 912)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1700  row: i  column: 06
            High quality sequence stop: 889.
FEATURES             Location/Qualifiers
     source          1. .912
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:4853837"
                     /tissue_type="primary B-cells from tonsils (cell line)"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_48"
                     /note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
                     Site_2: EcoRI; cDNA made by oligo-dT priming.
                     Directionally cloned into EcoRI/XhoI sites using the
                     following 5' adaptor: GGCACGAG(G). Size-selected >500bp
                     for average insert size 1.8kb. Library constructed by Ling
                     Hong in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using  ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match            80.4%;  Score 272.6;  DB 4;  Length 912;
  Best Local Similarity  95.3%;  Pred. No. 3.8e-71;
  Matches  281;  Conservative    0;  Mismatches   14;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db        121 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 180

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              ||||||||||||||||||||||||    |  ||||| |||| ||||||||||||||||||||
Db        181 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGTTATGGCATGAACTGGGTCCGCCAGGCTC 240

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db        241 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAATAATAAATACTATG 300

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 360

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA 296
              |||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        361 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGA 415


RESULT 9
BX344075
LOCUS       BX344075                 413 bp    mRNA    linear   EST 08-APR-2004
DEFINITION  BX344075 Homo sapiens PLACENTA COT 25-NORMALIZED Homo sapiens cDNA
            clone CS0DI052YB21 5-PRIME, mRNA sequence.
ACCESSION   BX344075
VERSION     BX344075.2  GI:46279625
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 413)
  AUTHORS   Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
  TITLE     Full-length cDNA libraries and normalization
  JOURNAL   Unpublished (2001)
COMMENT     On May 2, 2003 this sequence version replaced gi:30342243.
            Contact: Genoscope
            Genoscope - Centre National de Sequencage
            2 rue Gaston Cremieux, CP 5706 - 91057 EVRY cedex - FRANCE
            Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr
            1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime
            end enriched, double-strand cDNA was digested with Not I and cloned
            into the Not I and EcoR V sites of the pCMVSPORT 6 vector. Library
            was normalized. Library was constructed by Life Technologies, a
            division of Invitrogen. This sequence belongs to sequence cluster
            7198.r
            For more information about this cluster, see
            http://www.genoscope.cns.fr/cdna?s=CS1AI013ZH11QP1&c=7198.r.
FEATURES             Location/Qualifiers
     source          1. .413
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="CS0DI052YB21"
                     /tissue_type="PLACENTA COT 25-NORMALIZED"
                     /clone_lib="Homo sapiens PLACENTA COT 25-NORMALIZED"
                     /note="1st strand cDNA was primed with a NotI-oligo(dT)
                     primer. Five prime end enriched, double-strand cDNA was
                     digested with Not I and cloned into the Not I and EcoR V
                     sites of the pCMVSPORT 6 vector. Library was normalized."
ORIGIN

  Query Match            80.2%;  Score 271.8;  DB 5;  Length 413;
  Best Local Similarity  95.2%;  Pred. No. 5.6e-71;
  Matches  279;  Conservative    1;  Mismatches   13;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db        120 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 179

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              ||||||||||||||||||||||||    |   |||| ||||||||||||||||||||||||
Db        180 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 239
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Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||||||||||||||||||||||||||||:|||||||  ||||
Db        240 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTWATAAATATTATG 299

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        300 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 359

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAA 294
              ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        360 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAA 412


RESULT 10
AW401728
LOCUS       AW401728                516 bp    mRNA    linear   EST 16-FEB-2000
DEFINITION  UI-HF-BK0-aaf-f-12-0-UI.r1 NIH_MGC_36 Homo sapiens cDNA clone
            IMAGE:3053711 5', mRNA sequence.
ACCESSION   AW401728
VERSION     AW401728.1  GI:6920414
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 516)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Eco RI site shown at the beginning of the sequence.
             Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: M.B. Soares Lab
             cDNA Library Arrayed by: M.B. Soares Lab
             DNA Sequencing by: M.B. Soares Lab
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            www-bio.llnl.gov/bbrp/image/image.html
            Seq primer: M13 Forward.
FEATURES             Location/Qualifiers
     source          1. .516
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:3053711"
                     /tissue_type="lymph"
                     /cell_type="germinal center B cells"
                     /cell_line="MGC85"
                     /lab_host="DH10B (LTI)"
                     /clone_lib="NIH_MGC_36"
                     /note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI;
                     Constructed from size fractionated cytoplasmic mRNA
                     (0.5-1.5kb). Directionally cloned. Cells provided by Louis
                     M. Staudt, Ph.D. Library preparation by Maria de Fatima
                     Bonaldo, Ph.D.  and M. Bento Soares, Ph.D."
ORIGIN

  Query Match             79.8%;  Score 270.6;  DB 2;  Length 516;
  Best Local Similarity   95.2%;  Pred. No. 1.4e-70;
  Matches  279;  Conservative    0;  Mismatches   14;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||||
Db        132 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 191

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||    |   ||||| ||||||||||||||||||||||||
Db        192 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGCTATGCACTGGGTCCGCCAGGCTC 251

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        252 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTACG 311

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        312 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 371

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAA 294
              |||||||||||||||||||| ||||||||||||||||||||||||||||||
Db        372 TGCAAATGAACAGCCTGAGAACTGAGGACACGGCTGTGTATTACTGTGCGAAA 424


RESULT 11
BQ420418
LOCUS       BQ420418                899 bp    mRNA    linear   EST 23-MAY-2002
DEFINITION  AGENCOURT_7827143 NIH_MGC_92 Homo sapiens cDNA clone IMAGE:6014112
            5', mRNA sequence.
ACCESSION   BQ420418
VERSION     BQ420418.1  GI:21115733
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 899)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: ATCC
             cDNA Library Preparation: Life Technologies, Inc.
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Agencourt Bioscience Corporation
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLAM13208  row: b  column: 01
            High quality sequence stop: 500.
FEATURES             Location/Qualifiers
     source          1. .899
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:6014112"
                     /tissue_type="embryonal carcinoma, cell line"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_92"
                     /note="Organ: testis; Vector: pCMV-SPORT6; Site_1: NotI;
                     Site_2: SalI; Cloned unidirectionally; oligo-dT primed.
                     Average insert size 2.5 kb. Library enriched for
                     full-length clones and constructed by Life Technologies.
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             79.5%;  Score 269.6;  DB 5;  Length 899;
  Best Local Similarity   93.7%;  Pred. No. 3.1e-70;
  Matches  281;  Conservative    0;  Mismatches   19;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||||
Db         78 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 137

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||    |   ||||| ||||||||||||||||||||||||
Db        138 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGCTATGCACTGGGTCCGCCAGGCTC 197

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |||||||||| |
Db        198 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCAATAAATACTACG 257

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        258 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 317
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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              ||||||||||||||||||||||||||||||||||||||||||||||||||| | |   ||
Db        318 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAGAGACACAG 377


RESULT 12
BG686759
LOCUS       BG686759                663 bp    mRNA    linear   EST 01-MAY-2001
DEFINITION  602650729F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4763215 5',
            mRNA sequence.
ACCESSION   BG686759
VERSION     BG686759.1  GI:13918156
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 663)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1618  row: i  column: 08
            High quality sequence stop: 659.
FEATURES             Location/Qualifiers
     source          1. .663
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:4763215"
                     /tissue_type="primary B-cells from tonsils (cell line)"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_48"
                     /note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
                     Site_2: EcoRI; cDNA made by oligo-dT priming.
                     Directionally cloned into EcoRI/XhoI sites using the
                     following 5' adaptor: GGCACGAG(G). Size-selected >500bp
                     for average insert size 1.8kb. Library constructed by Ling
                     Hong in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using  ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             79.5%;  Score 269.4;  DB 4;  Length 663;
  Best Local Similarity   94.6%;  Pred. No. 3.3e-70;
  Matches  279;  Conservative    0;  Mismatches   16;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Db        129 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 188

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Db        189 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGCTATGCACTGGGTCCGCCAGGCTC 248

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        249 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCAATAAATACTACG 308

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Db        309 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 368

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAA 296
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Db        369 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAGAGA 423


RESULT 13
AW402572
LOCUS       AW402572                503 bp    mRNA    linear   EST 16-FEB-2000
DEFINITION  UI-HF-BK0-aax-b-12-0-UI.r1 NIH_MGC_36 Homo sapiens cDNA clone
            IMAGE:3055079 5', mRNA sequence.
ACCESSION   AW402572
VERSION     AW402572.1  GI:6921271
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 503)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Eco RI site shown at the beginning of the sequence.
             Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: M.B. Soares Lab
             cDNA Library Arrayed by: M.B. Soares Lab
             DNA Sequencing by: M.B. Soares Lab
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            www-bio.llnl.gov/bbrp/image/image.html
            Seq primer: M13 Forward.
FEATURES             Location/Qualifiers
     source          1. .503
                     /organism="Homo sapiens"
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                     /clone="IMAGE:3055079"
                     /tissue_type="lymph"
                     /cell_type="germinal center B cells"
                     /cell_line="MGC85"
                     /lab_host="DH10B (LTI)"
                     /clone_lib="NIH_MGC_36"
                     /note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI;
                     Constructed from size fractionated cytoplasmic mRNA
                     (0.5-1.5kb). Directionally cloned. Cells provided by Louis
                     M. Staudt, Ph.D. Library preparation by Maria de Fatima
                     Bonaldo, Ph.D.  and M. Bento Soares, Ph.D."
ORIGIN

  Query Match             78.8%;  Score 267.2;  DB 2;  Length 503;
  Best Local Similarity   87.0%;  Pred. No. 1.4e-69;
  Matches  315;  Conservative    0;  Mismatches   23;  Indels   24;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
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Db         44 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 103

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Db        104 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGCTATGCACTGGGTCCGCCAGGCTC 163

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAATAATAAATATTACG 181
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Db        164 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAATAATAAATATTACG 223

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Db        224 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 283

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAA------ 295
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Qy        296 ------------------AGGAAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCT 337
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Db        344 ATTACGATTTTTGGAGTGCCACAAACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCT 403

Qy        338 CA 339
              ||
Db        404 CA 405


RESULT 14
BQ708235
LOCUS       BQ708235                894 bp    mRNA    linear   EST 16-JUL-2002
DEFINITION  AGENCOURT_8354494 NIH_MGC_113 Homo sapiens cDNA clone IMAGE:6281304
            5', mRNA sequence.
ACCESSION   BQ708235
VERSION     BQ708235.1  GI:21847134
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 894)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Dr. Mark Watson
             cDNA Library Preparation: Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Agencourt Bioscience Corporation
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM2473  row: o  column: 01
            High quality sequence stop: 585.
FEATURES             Location/Qualifiers
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                     /clone="IMAGE:6281304"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_113"
                     /note="Organ: spleen; Vector: pOTB7; Site_1: XhoI; Site_2:
                     EcoRI; cDNA made by oligo-dT priming. Directionally cloned
                     into EcoRI/XhoI sites using the following 5' adaptor:
                     GGCACGAG(G). Library constructed  by Ling Hong in the
                     laboratory of Gerald M. Rubin (University of  California,
                     Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and
                     Superscript II RT (Life Technologies). Note: this is a
                     NIH_MGC Library."
ORIGIN

  Query Match             78.7%;  Score 266.8;  DB 5;  Length 894;
  Best Local Similarity   92.7%;  Pred. No. 2.2e-69;
  Matches  280;  Conservative    0;  Mismatches   22;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | ||||||||||||| |||||| ||||||||||||||||||||||||||
Db        122 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGGGGTCCCTGAGACTCT 181

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Db        182 CCTGTGCAGCGTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 241

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        242 CAGGGAAGGGGCTGGAGTGGGTGGCATTTATACGATATGATGGAAGTAATAAATACTACG 301

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              |||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||| | |
Db        362 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAATTAAAGG 421

Qy        302 GC 303
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Db        422 GC 423


RESULT 15
BG503730
LOCUS       BG503730                456 bp    mRNA    linear   EST 27-MAR-2001
DEFINITION  602549705F1 NIH_MGC_61 Homo sapiens cDNA clone IMAGE:4657248 5',
            mRNA sequence.
ACCESSION   BG503730
VERSION     BG503730.1  GI:13465247
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 456)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: ATCC
             cDNA Library Preparation: CLONETECH Laboratories, Inc.
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1449  row: j  column: 01
            High quality sequence stop: 456.
FEATURES             Location/Qualifiers
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                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:4657248"
                     /tissue_type="embryonal carcinoma"
                     /lab_host="DH10B (T1 phage-resistant)"
                     /clone_lib="NIH_MGC_61"
                     /note="Organ: testis; Vector: pDNR-LIB (Clontech); Site_1:
                     SfiI (ggccgcctcggcc); Site_2: SfiI (ggccattatggcc);
                     Double-stranded cDNA was prepared from cell line RNA.  5'
                     and 3' adaptors were used in cloning as follows: 5'
                     adaptor sequence:  5'-CACGGCCATTATGGCC-3' and 3' adaptor
                     sequence:  5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3'
                     (where B = A, C, or G and N = A, C,  G, or T).  Average
                     insert size 1.75 kb (range 0.9-4.0 kb). 15/15 colonies
                     contained  inserts by PCR. This library was enriched for
                     full-length clones and was constructed by Clontech
                     Laboratories (Palo Alto, CA). Note: this is a NIH_MGC
                     Library."
ORIGIN

  Query Match             78.6%;  Score 266.6;  DB 4;  Length 456;
  Best Local Similarity   90.7%;  Pred. No. 2.1e-69;
  Matches  284;  Conservative    0;  Mismatches   29;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db         96 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 155

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| ||||||||||||||    |    |||| ||||||||||||||||||||||||
Db        156 CCTGTGCAGCGTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 215

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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                ||||||||||||||||||||||||||||||||  ||||||||||||||||||||||||
Db        216 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATGGTATGATGGAAGTAATAAATACTATG 275

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        276 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 335

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
                |||||||||||||||||||| |||||||||||||||||||||||| | |    ||
Db        336 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGACACAG 395

Qy        302 GCTACTGGGGCCA 314
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Db        396 TGAGGGGAGGTCA 408
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

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Searched: 1202784 seqs, 818138359 residues

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	274.4	80.9	675	4	US-09-456-090A-67	Sequence 67, Appl
2	274.4	80.9	675	4	US-09-453-234-67	Sequence 67, Appl
3	273.6	80.7	369	4	US-09-560-198A-9	Sequence 9, Appli
4	272.8	80.5	512	3	US-08-545-809A-30	Sequence 30, Appl
5	272.8	80.5	675	4	US-09-456-090A-59	Sequence 59, Appl
6	272.8	80.5	675	4	US-09-456-090A-91	Sequence 91, Appl
7	272.8	80.5	675	4	US-09-453-234-59	Sequence 59, Appl
8	272.8	80.5	675	4	US-09-453-234-91	Sequence 91, Appl
9	272.6	80.4	369	4	US-09-560-198A-1	Sequence 1, Appli
10	272.2	80.3	677	4	US-09-456-090A-55	Sequence 55, Appl
11	272.2	80.3	677	4	US-09-453-234-55	Sequence 55, Appl
12	272	80.2	369	4	US-09-560-198A-3	Sequence 3, Appli
13	271.6	80.1	369	4	US-09-424-840B-5	Sequence 5, Appli
14	271.4	80.1	1392	4	US-09-472-087-30	Sequence 30, Appl
15	271.4	80.1	1392	4	US-09-472-087-59	Sequence 59, Appl
16	271.2	80.0	675	4	US-09-456-090A-101	Sequence 101, App
17	271.2	80.0	675	4	US-09-453-234-101	Sequence 101, App
18	270	79.6	1395	4	US-09-472-087-28	Sequence 28, Appl
19	270	79.6	1395	4	US-09-472-087-57	Sequence 57, Appl
20	269.8	79.6	1392	4	US-09-472-087-27	Sequence 27, Appl
21	269.8	79.6	1392	4	US-09-472-087-53	Sequence 53, Appl
22	269.8	79.6	1392	4	US-09-472-087-55	Sequence 55, Appl
23	269.8	79.6	1999	4	US-09-472-087-54	Sequence 54, Appl
24	268	79.1	675	4	US-09-456-090A-107	Sequence 107, App
25	268	79.1	675	4	US-09-453-234-107	Sequence 107, App
26	266.4	78.6	675	4	US-09-456-090A-99	Sequence 99, Appl
27	266.4	78.6	675	4	US-09-456-090A-105	Sequence 105, App
28	266.4	78.6	675	4	US-09-456-090A-109	Sequence 109, App
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30	266.4	78.6	675	4	US-09-453-234-105	Sequence 105, App
31	266.4	78.6	675	4	US-09-453-234-109	Sequence 109, App
32	266.2	78.5	1413	4	US-09-472-087-61	Sequence 61, Appl
33	265.8	78.4	675	4	US-09-456-090A-93	Sequence 93, Appl
34	265.8	78.4	675	4	US-09-453-234-93	Sequence 93, Appl
35	265.8	78.4	5925	4	US-09-315-926A-78	Sequence 78, Appl
36	264.2	77.9	372	3	US-09-240-274-73	Sequence 73, Appl
37	264.2	77.9	372	3	US-09-240-274-74	Sequence 74, Appl
38	264.2	77.9	372	3	US-09-240-274-75	Sequence 75, Appl
39	263.8	77.8	372	4	US-09-424-840B-15	Sequence 15, Appl
40	263.8	77.8	375	3	US-09-240-274-89	Sequence 89, Appl
41	263.8	77.8	384	3	US-09-240-274-70	Sequence 70, Appl
42	263.6	77.8	423	1	US-08-259-372A-1	Sequence 1, Appli
43	263.6	77.8	423	1	US-08-468-671-1	Sequence 1, Appli
44	263.2	77.6	675	4	US-09-456-090A-57	Sequence 57, Appl
45	263.2	77.6	675	4	US-09-456-090A-61	Sequence 61, Appl

ALIGNMENTS

RESULT 1
US-09-456-090A-67
; Sequence 67, Application US/09456090A
; Patent No. 6680209
; GENERAL INFORMATION:
; APPLICANT: Buechler, Joe
; APPLICANT: Valkirs, Gunars
; APPLICANT: Gray, Jeff
; APPLICANT: Lonberg, Nils
; TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
; FILE REFERENCE: 020015-000200US
; CURRENT APPLICATION NUMBER: US/09/456,090A
; CURRENT FILING DATE: 1999-12-06
; NUMBER OF SEQ ID NOS: 110
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 67
; LENGTH: 675
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)..(675)
; OTHER INFORMATION: M1-23H
US-09-456-090A-67

Query Match 80.9%; Score 274.4; DB 4; Length 675;
Best Local Similarity 89.1%; Pred. No. 1.9e-80;
Matches 312; Conservative 0; Mismatches 26; Indels 12; Gaps 1;

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Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              ||||||||||| |  |||||||||||| |||||| |||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGCAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| |||||||||||||    | | |||| |||||||||||||||||||||||||
Db         62 CCTGTGCAGCGTCTGGATTCACCTTCAGTAACTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGCTATATGGTATGATGGAAGTAAAACATACAATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
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RESULT 2
US-09-453-234-67
; Sequence 67, Application US/09453234
; Patent No. 6794132
; GENERAL INFORMATION:
;  APPLICANT: Buechler, Joe
;  APPLICANT:  Valkirs, Gunars
;  APPLICANT:  Gray, Jeff
;  APPLICANT:  Lonberg, Nils
;  APPLICANT:  Biosite Diagnostics, Inc.
;  APPLICANT:  GenPharm International
;  TITLE OF INVENTION: Human Antibodies
;  FILE REFERENCE: 020015-000110US
;  CURRENT APPLICATION NUMBER: US/09/453,234
;  CURRENT FILING DATE:  1999-12-01
;  PRIOR APPLICATION NUMBER: US 60/157,415
;  PRIOR FILING DATE: 1999-10-02
;  NUMBER OF SEQ ID NOS: 112
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 67
;   LENGTH: 675
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: M1-23H
;   NAME/KEY: CDS
;   LOCATION: (1)..(675)
US-09-453-234-67

  Query Match             80.9%;  Score 274.4;  DB 4;  Length 675;
  Best Local Similarity   89.1%;  Pred. No. 1.9e-80;
  Matches  312;  Conservative    0;  Mismatches   26;  Indels   12;  Gaps    1;

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              |||||||||||| |  |||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGCAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
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Db         62 CCTGTGCAGCGTCTGGATTCACCTTCAGTAACTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||| |||||  ||||||||||||| | |||| |||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGCTATATGGTATGATGGAAGTAAAACATACAATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||||||||||| |||||||||||||||||||||||||| | | |
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGATGGGA 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | ||||||||||||||||||||||||||||||||||||||||
Db        302 TAGGCTACTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351


RESULT 3
US-09-560-198A-9
; Sequence 9, Application US/09560198A
; Patent No. 6492497
; GENERAL INFORMATION:
;  APPLICANT: Thompson, Julia E
;  APPLICANT:  Lennard, Simon N
;  APPLICANT:  Wilton, Alison J
;  APPLICANT:  Braddock, Peta SH
;  APPLICANT:  Du Fou, Sarah L
;  APPLICANT:  McCafferty, John G
;  APPLICANT:  Conroy, Louise A
;  APPLICANT:  Tempest, Philip R
;  TITLE OF INVENTION: Specific binding members for TGFbeta1
;  FILE REFERENCE: 28111/35620A
;  CURRENT APPLICATION NUMBER: US/09/560,198A
;  CURRENT FILING DATE:  2000-04-28
;  PRIOR APPLICATION NUMBER: US 60/131,983
;  PRIOR FILING DATE: 1999-04-30
;  NUMBER OF SEQ ID NOS: 25
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 9
;   LENGTH: 369
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-560-198A-9

  Query Match             80.7%;  Score 273.6;  DB 4;  Length 369;
  Best Local Similarity   89.6%;  Pred. No. 2.6e-80;
  Matches  294;  Conservative    0;  Mismatches   34;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||| ||||||||||||||||||||||||||||||||||||||| ||||||||||
Db        122 CAGGCAAGGAGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTATTAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              |||||||||||||||||||||||||||||||||||||||||||||||||||||  ||  | |
Db        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGCGAACTGGTG 301

Qy        302 GCTACTGGGGCCAGGGAACCCTGGTCAC 329
                ||  | ||| | |  || |  | | |
Db        302 AATATAGTGGCTACGATACGCCCGCCTC 329


RESULT 4
US-08-545-809A-30
; Sequence 30, Application US/08545809A
; Patent No. 6096878
;  GENERAL INFORMATION:
;    APPLICANT:  Honjo, Tasuku
;    APPLICANT:  Matsuda, Fumihiko
;    TITLE OF INVENTION:  HUMAN IMMUNOGLOBULIN VH GENE
;    TITLE OF INVENTION:  SEGMENTS AND DNA FRAGMENTS CONTAINING THE SAME
;    NUMBER OF SEQUENCES:  145
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Fish & Richardson, P.C.
;      STREET:  225 Franklin Street
;      CITY:  Boston
;      STATE:  MA
;      COUNTRY:  US
;      ZIP:  02110-2804
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Diskette
;      COMPUTER:  IBM Compatible
;      OPERATING SYSTEM:  Windows95
;      SOFTWARE:  FastSEQ for Windows Version 2.0
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/545,809A
;      FILING DATE:  27-MAR-1996
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  PCT/JP93/00603
;      FILING DATE:  10-MAY-1993
```

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;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Freeman, John W.
;      REGISTRATION NUMBER:  29,066
;      REFERENCE/DOCKET NUMBER:  06501/004001
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  617-542-5070
;      TELEFAX:  617-542-8906
;      TELEX:  200154
;  INFORMATION FOR SEQ ID NO:  30:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  512 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  Genomic DNA
;    ORIGINAL SOURCE:
;      ORGANISM:  Homo sapiens
;      CELL TYPE:  human lymphoblast
;      CELL LINE:  CGM1
US-08-545-809A-30

  Query Match            80.5%;  Score 272.8;  DB 3;  Length 512;
  Best Local Similarity  94.3%;  Pred. No. 5.6e-80;
  Matches  283;  Conservative    0;  Mismatches   17;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db        169 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 228

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||||
Db        229 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 288

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        289 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 348

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        349 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 408

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              |||||||||||||||||||||||||||||||||||||||||||||||||||| | |   ||
Db        409 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAGAGACACAG 468


RESULT 5
US-09-456-090A-59
; Sequence 59, Application US/09456090A
; Patent No. 6680209
; GENERAL INFORMATION:
;  APPLICANT: Buechler, Joe
;  APPLICANT:  Valkirs, Gunars
;  APPLICANT:  Gray, Jeff
;  APPLICANT:  Lonberg, Nils
;  TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
;  FILE REFERENCE: 020015-000200US
;  CURRENT APPLICATION NUMBER: US/09/456,090A
;  CURRENT FILING DATE:  1999-12-06
;  NUMBER OF SEQ ID NOS: 110
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 59
;   LENGTH: 675
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(675)
;   OTHER INFORMATION: M1-5H
US-09-456-090A-59

  Query Match            80.5%;  Score 272.8;  DB 4;  Length 675;
  Best Local Similarity  88.9%;  Pred. No. 6.4e-80;
  Matches  311;  Conservative    0;  Mismatches   27;  Indels   12;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| |||||||| |||||    | | |||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCGTCTGGATTTACCTTCAGTTACTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||| || ||||| | ||||||||  ||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGACACTTATAACCTATGATGGAGATAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||||||||||||||||||||||||||||||||||||||||||| | | |
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGACGGGA 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | ||| |||||||||||||||||||||||||||||||||
Db        302 TCGGGTACTTTGACTATTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351


RESULT 6
US-09-456-090A-91
; Sequence 91, Application US/09456090A
; Patent No. 6680209
; GENERAL INFORMATION:
;  APPLICANT: Buechler, Joe
;  APPLICANT:  Valkirs, Gunars
;  APPLICANT:  Gray, Jeff
;  APPLICANT:  Lonberg, Nils
;  TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
;  FILE REFERENCE: 020015-000200US
;  CURRENT APPLICATION NUMBER: US/09/456,090A
;  CURRENT FILING DATE:  1999-12-06
;  NUMBER OF SEQ ID NOS: 110
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 91
;   LENGTH: 675
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(675)
;   OTHER INFORMATION: M2-11H
US-09-456-090A-91

  Query Match            80.5%;  Score 272.8;  DB 4;  Length 675;
  Best Local Similarity  88.9%;  Pred. No. 6.4e-80;
  Matches  311;  Conservative    0;  Mismatches   27;  Indels   12;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| |||||||| |||||    | | |||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCGTCTGGATTTACCTTCAGTTACTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              ||||||||||||||||||||||| || ||||| | ||||||||  ||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGACACTTATAACCTATGATGGAGATAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
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                ||||||||||||||||||||| |||||||||||||||||||||||||||||| | | |
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGACGGGA 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | ||| ||||||||||||||||||||||||||||||||||||
Db        302 TCGGGTACTTTGACTATTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351


RESULT 7
US-09-453-234-59
; Sequence 59, Application US/09453234
; Patent No. 6794132
; GENERAL INFORMATION:
;  APPLICANT: Buechler, Joe
;  APPLICANT:  Valkirs, Gunars
;  APPLICANT:  Gray, Jeff
;  APPLICANT:  Lonberg, Nils
;  APPLICANT:  Biosite Diagnostics, Inc.
;  APPLICANT:  GenPharm International
;  TITLE OF INVENTION: Human Antibodies
;  FILE REFERENCE: 020015-000110US
;  CURRENT APPLICATION NUMBER: US/09/453,234
;  CURRENT FILING DATE:  1999-12-01
;  PRIOR APPLICATION NUMBER: US 60/157,415
;  PRIOR FILING DATE: 1999-10-02
;  NUMBER OF SEQ ID NOS: 112
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 59
;   LENGTH: 675
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: M1-5H
;   NAME/KEY: CDS
;   LOCATION: (1)..(675)
US-09-453-234-59

  Query Match             80.5%;  Score 272.8;  DB 4;  Length 675;
  Best Local Similarity   88.9%;  Pred. No. 6.4e-80;
  Matches  311;  Conservative    0;  Mismatches   27;  Indels   12;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| |||||||| ||||    | | |||| ||||||||||||||||||||||
Db         62 CCTGTGCAGCGTCTGGATTTACCTTCAGTTACTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||| || ||||| | |||||||||  |||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGACACTTATAACCTATGATGGAGATAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||||||||||| |||||||||||||||||||||||||||| | | |
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGACGGGA 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | ||| ||||||||||||||||||||||||||||||||||||
Db        302 TCGGGTACTTTGACTATTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351


RESULT 8
US-09-453-234-91
; Sequence 91, Application US/09453234
; Patent No. 6794132
; GENERAL INFORMATION:
;  APPLICANT: Buechler, Joe
;  APPLICANT:  Valkirs, Gunars
;  APPLICANT:  Gray, Jeff
;  APPLICANT:  Lonberg, Nils
;  APPLICANT:  Biosite Diagnostics, Inc.
;  APPLICANT:  GenPharm International
;  TITLE OF INVENTION: Human Antibodies
;  FILE REFERENCE: 020015-000110US
;  CURRENT APPLICATION NUMBER: US/09/453,234
;  CURRENT FILING DATE:  1999-12-01
;  PRIOR APPLICATION NUMBER: US 60/157,415
;  PRIOR FILING DATE: 1999-10-02
;  NUMBER OF SEQ ID NOS: 112
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 91
;   LENGTH: 675
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(675)
;   OTHER INFORMATION: M2-11H
US-09-453-234-91

  Query Match             80.5%;  Score 272.8;  DB 4;  Length 675;
  Best Local Similarity   88.9%;  Pred. No. 6.4e-80;
  Matches  311;  Conservative    0;  Mismatches   27;  Indels   12;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| |||||||| ||||    | | |||| ||||||||||||||||||||||
Db         62 CCTGTGCAGCGTCTGGATTTACCTTCAGTTACTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||| || ||||| | |||||||||  |||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGACACTTATAACCTATGATGGAGATAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              |||||||||||||||||||| |||||||||||||||||||||||||||| | | |
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGACGGGA 301

Qy        299 ---------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      | ||| ||||||||||||||||||||||||||||||||||||
Db        302 TCGGGTACTTTGACTATTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351


RESULT 9
US-09-560-198A-1
; Sequence 1, Application US/09560198A
; Patent No. 6492497
; GENERAL INFORMATION:
;  APPLICANT: Thompson, Julia E
;  APPLICANT:  Lennard, Simon N
;  APPLICANT:  Wilton, Alison J
;  APPLICANT:  Braddock, Peta SH
;  APPLICANT:  Du Fou, Sarah L
;  APPLICANT:  McCafferty, John G
;  APPLICANT:  Conroy, Louise A
;  APPLICANT:  Tempest, Philip R
;  TITLE OF INVENTION: Specific binding members for TGFbeta1
;  FILE REFERENCE: 28111/35620A
;  CURRENT APPLICATION NUMBER: US/09/560,198A
;  CURRENT FILING DATE:  2000-04-28
;  PRIOR APPLICATION NUMBER: US 60/131,983
;  PRIOR FILING DATE: 1999-04-30
;  NUMBER OF SEQ ID NOS: 25
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
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;   LENGTH: 369
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-560-198A-1

  Query Match             80.4%;  Score 272.6;  DB 4;  Length 369;
  Best Local Similarity   90.9%;  Pred. No. 5.5e-80;
  Matches  290;  Conservative    0;  Mismatches   29;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||  |  ||||||||||| ||||||  |||||||||  ||||||||||||||||
Db          2 AGGTGCAGCTGGTGCAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTATTAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              |||||||||||||||||||||||||||||||||||||||||||||||||  ||  |  |
Db        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGCGAACTGGTG 301

Qy        302 GCTACTGGGGCCAGGGAAC 320
                ||  | ||| | |  ||
Db        302 AATATAGTGGCTACGATAC 320


RESULT 10
US-09-456-090A-55
; Sequence 55, Application US/09456090A
; Patent No. 6680209
; GENERAL INFORMATION:
;  APPLICANT: Buechler, Joe
;  APPLICANT:  Valkirs, Gunars
;  APPLICANT:  Gray, Jeff
;  APPLICANT:  Lonberg, Nils
;  TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
;  FILE REFERENCE: 020015-000200US
;  CURRENT APPLICATION NUMBER: US/09/456,090A
;  CURRENT FILING DATE:  1999-12-06
;  NUMBER OF SEQ ID NOS: 110
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 55
;   LENGTH: 677
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (3)..(677)
;   OTHER INFORMATION: M1-3H
US-09-456-090A-55

  Query Match             80.3%;  Score 272.2;  DB 4;  Length 677;
  Best Local Similarity   88.6%;  Pred. No. 1e-79;
  Matches  311;  Conservative    0;  Mismatches   28;  Indels   12;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              || |||||||||  |  ||||||||||| ||||||  |||||||||  ||||||||||||||
Db          3 GATGTGCAGCTGGTGCAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 62

Qy         61 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              |||||||||||  |||||||||||||||    | | |||| ||||||||||||||||||||||
Db         63 TCCTGTGCAGCGTCTGGATTCACCTTCAGTTACTATGGCATGCACTGGGTCCGCCAGGCT 122

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
              |||||||||||||||||||||||| || ||||| | |||||||||  ||||||||||||||
Db        123 CCAGGCAAGGGGCTGGAGTGGGTGACACTTATAACCTATGATGGAGATAATAAATACTAT 182

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        183 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 242

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG-- 298
              |||||||||||||||||||||||| |||||||||||||||||||||||||||||| | | |
Db        243 CTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGACGGG 302

Qy        299 ----------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                         | ||| ||||||||||||||||||||||||||||||||
Db        303 ATCGGGTACTTTGACTATTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 353


RESULT 11
US-09-453-234-55
; Sequence 55, Application US/09453234
; Patent No. 6794132
; GENERAL INFORMATION:
;  APPLICANT: Buechler, Joe
;  APPLICANT:  Valkirs, Gunars
;  APPLICANT:  Gray, Jeff
;  APPLICANT:  Lonberg, Nils
;  APPLICANT:  Biosite Diagnostics, Inc.
;  APPLICANT:  GenPharm International
;  TITLE OF INVENTION: Human Antibodies
;  FILE REFERENCE: 020015-000110US
;  CURRENT APPLICATION NUMBER: US/09/453,234
;  CURRENT FILING DATE:  1999-12-01
;  PRIOR APPLICATION NUMBER: US 60/157,415
;  PRIOR FILING DATE: 1999-10-02
;  NUMBER OF SEQ ID NOS: 112
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 55
;   LENGTH: 677
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: M1-3H
;   NAME/KEY: CDS
;   LOCATION: (3)..(677)
US-09-453-234-55

  Query Match             80.3%;  Score 272.2;  DB 4;  Length 677;
  Best Local Similarity   88.6%;  Pred. No. 1e-79;
  Matches  311;  Conservative    0;  Mismatches   28;  Indels   12;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              || |||||||||  |  ||||||||||| ||||||  |||||||||  ||||||||||||||
Db          3 GATGTGCAGCTGGTGCAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 62

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              |||||||||||  |||||||||||||||    | | |||| ||||||||||||||||||||||
Db         63 TCCTGTGCAGCGTCTGGATTCACCTTCAGTTACTATGGCATGCACTGGGTCCGCCAGGCT 122

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
              |||||||||||||||||||||||| || ||||| | |||||||||  ||||||||||||||
Db        123 CCAGGCAAGGGGCTGGAGTGGGTGACACTTATAACCTATGATGGAGATAATAAATACTAT 182

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        183 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 242

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG-- 298
              |||||||||||||||||||||||| |||||||||||||||||||||||||||||| | | |
Db        243 CTGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGACGGG 302

Qy        299 ----------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                         | ||| ||||||||||||||||||||||||||||||||
Db        303 ATCGGGTACTTTGACTATTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 353
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RESULT 12
US-09-560-198A-3
; Sequence 3, Application US/09560198A
; Patent No. 6492497
; GENERAL INFORMATION:
;  APPLICANT: Thompson, Julia E
;  APPLICANT:  Lennard, Simon N
;  APPLICANT:  Wilton, Alison J
;  APPLICANT:  Braddock, Peta SH
;  APPLICANT:  Du Fou, Sarah L
;  APPLICANT:  McCafferty, John G
;  APPLICANT:  Conroy, Louise A
;  APPLICANT:  Tempest, Philip R
;  TITLE OF INVENTION: Specific binding members for TGFbeta1
;  FILE REFERENCE: 28111/35620A
;  CURRENT APPLICATION NUMBER: US/09/560,198A
;  CURRENT FILING DATE:  2000-04-28
;  PRIOR APPLICATION NUMBER: US 60/131,983
;  PRIOR FILING DATE: 1999-04-30
;  NUMBER OF SEQ ID NOS: 25
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
;   LENGTH: 369
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-560-198A-3

  Query Match             80.2%;  Score 272;  DB 4;  Length 369;
  Best Local Similarity   90.6%;  Pred. No. 8.7e-80;
  Matches  290;  Conservative    0;  Mismatches   30;  Indels    0;  Gaps    0;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              ||||| |||||| | ||||||||||||| |||||| ||||||||| ||||||||||||||
Db          1 GAGGTCCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 60

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              |||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         61 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 120

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
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Db        121 CCAGGCAAGGAGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTATTAAATACTAT 180

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
              |||||||||||||||||||||||||||||||||||||||||||||||||||  ||  |
Db        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGCGAACTGGT 300

Qy        301 GGCTACTGGGGCCAGGGAAC 320
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Db        301 GAATATAGTGGCTACGATAC 320


RESULT 13
US-09-424-840B-5
; Sequence 5, Application US/09424840B
; Patent No. 6790938
; GENERAL INFORMATION:
;  APPLICANT: Berchtold, Peter
;  APPLICANT:  Escher, Robert F. A.
;  TITLE OF INVENTION: ANTI-GPIIB/IIIA RECOMBINANT ANTIBODIES
;  FILE REFERENCE: 100564-09049
;  CURRENT APPLICATION NUMBER: US/09/424,840B
;  CURRENT FILING DATE:  1999-12-03
;  PRIOR APPLICATION NUMBER: DE 19820663.1
;  PRIOR FILING DATE: 1998-05-08
;  PRIOR APPLICATION NUMBER: DE 19755227.7
;  PRIOR FILING DATE: 1997-12-12
;  PRIOR APPLICATION NUMBER: DE 19723904.8
;  PRIOR FILING DATE: 1997-06-06
;  NUMBER OF SEQ ID NOS: 128
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
;   LENGTH: 369
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(369)
;   OTHER INFORMATION:
US-09-424-840B-5

  Query Match             80.1%;  Score 271.6;  DB 4;  Length 369;
  Best Local Similarity   92.3%;  Pred. No. 1.2e-79;
  Matches  286;  Conservative    0;  Mismatches   24;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              ||||| | |||||||||||||||||| |||||| ||||||||| ||||||||||||||||
Db          2 AGGTGAAACTGCTCGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGCTATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCAATAAATACTACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||| |||||||||| |
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGCAATAAATACTACG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Db        182 CAGACTCCGTGAAGGGCCGATTCGCCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
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Db        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAGAGCGCTGG 301

Qy        302 GCTACTGGGG 311
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RESULT 14
US-09-472-087-30
; Sequence 30, Application US/09472087
; Patent No. 6682736
; GENERAL INFORMATION:
;  APPLICANT: HANSON, DOUGLAS C.
;  APPLICANT:  NEVEU, MARK J.
;  APPLICANT:  MUELLER, EILLEN E.
;  APPLICANT:  HANKE, JEFFREY H.
;  APPLICANT:  GILMAN, STEVEN C.
;  APPLICANT:  DAVIS, C. GEOFFREY
;  APPLICANT:  CORVALAN, JOSE R.
;  TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO CTLA-4
;  FILE REFERENCE: ABX-PF1
;  CURRENT APPLICATION NUMBER: US/09/472,087
;  CURRENT FILING DATE:  1999-12-23
;  PRIOR APPLICATION NUMBER: 60/113,647
;  PRIOR FILING DATE: 1998-12-23
;  NUMBER OF SEQ ID NOS: 147
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 30
;   LENGTH: 1392
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-472-087-30

  Query Match             80.1%;  Score 271.4;  DB 4;  Length 1392;
  Best Local Similarity   88.4%;  Pred. No. 2.6e-79;
  Matches  312;  Conservative    0;  Mismatches   26;  Indels   15;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| ||||||  |||||||| ||||||||||||||||
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Db         59 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCGAGCCTGGGAGGTCCCTGAGACTCT 118

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              ||||| |||| |||||||||||||||   | ||||| |||||||||||||||||||||||
Db        119 CCTGTACAGCGTCTGGATTCACCTTCAGTAGTTATGGCATGCACTGGGTCCGCCAGGCTC 178

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||  |||||||||| |||||| ||||||
Db        179 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATGGTATGATGGAAGCAATAAACACTATG 238

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
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Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA- 300
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Db        299 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGCCGGAC 358

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RESULT 15
US-09-472-087-59
; Sequence 59, Application US/09472087
; Patent No. 6682736
; GENERAL INFORMATION:
;  APPLICANT: HANSON, DOUGLAS C.
;  APPLICANT:  NEVEU, MARK J.
;  APPLICANT:  MUELLER, EILLEN E.
;  APPLICANT:  HANKE, JEFFREY H.
;  APPLICANT:  GILMAN, STEVEN C.
;  APPLICANT:  DAVIS, C. GEOFFREY
;  APPLICANT:  CORVALAN, JOSE R.
;  TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO CTLA-4
;  FILE REFERENCE: ABX-PF1
;  CURRENT APPLICATION NUMBER: US/09/472,087
;  CURRENT FILING DATE:  1999-12-23
;  PRIOR APPLICATION NUMBER: 60/113,647
;  PRIOR FILING DATE: 1998-12-23
;  NUMBER OF SEQ ID NOS: 147
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 59
;   LENGTH: 1392
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-472-087-59

  Query Match             80.1%;  Score 271.4;  DB 4;  Length 1392;
  Best Local Similarity   88.4%;  Pred. No. 2.6e-79;
  Matches  312;  Conservative    0;  Mismatches   26;  Indels   15;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | ||||||||||||| ||||||  |||||||| ||||||||||||||
Db         59 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCGAGCCTGGGAGGTCCCTGAGACTCT 118

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              ||||| |||| |||||||||||||||   | ||||| |||||||||||||||||||||||
Db        119 CCTGTACAGCGTCTGGATTCACCTTCAGTAGTTATGGCATGCACTGGGTCCGCCAGGCTC 178

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
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Db        179 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATGGTATGATGGAAGCAATAAACACTATG 238

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              |||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||
Db        239 CAGACTCCGCGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 298

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA- 300
              |||||||||||||||||||||| |||||||||||||||||||||||||||||| |   | |
Db        299 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGCCGGAC 358

Qy        301 --------------GGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
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Job time : 106.268 secs
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: September 11, 2005, 21:22:08 ; Search time 536.75 Seconds
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Perfect score: 339
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 7351250 seqs, 3283620254 residues

Total number of hits satisfying chosen parameters: 14702500

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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26: /cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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1	339	100.0	372	15	US-10-325-694-145	Sequence 145, App
2	301.2	88.8	729	18	US-10-406-830-20	Sequence 20, Appl
3	294.8	87.0	396	17	US-10-309-764-54	Sequence 54, Appl
4	289.2	85.3	342	20	US-10-738-120-9	Sequence 9, Appli
5	285.2	84.1	537	21	US-10-805-177-25	Sequence 25, Appl
6	284.4	83.9	357	16	US-10-010-729-8	Sequence 8, Appli
7	284	83.8	352	21	US-10-638-265-17	Sequence 17, Appl
8	283.8	83.7	405	17	US-10-309-764-58	Sequence 58, Appl
9	282	83.2	783	19	US-10-642-120-5	Sequence 5, Appli
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12	282	83.2	783	19	US-10-642-124-5	Sequence 5, Appli
13	282	83.2	783	19	US-10-621-269-5	Sequence 5, Appli
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26	282	83.2	783	22	US-10-642-059-5	Sequence 5, Appli
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28	280.6	82.8	349	19	US-10-684-109-18	Sequence 18, Appl
29	280.6	82.8	404	17	US-10-309-764-74	Sequence 74, Appl
30	280.6	82.8	405	17	US-10-309-764-70	Sequence 70, Appl
31	279	82.3	405	17	US-10-309-764-66	Sequence 66, Appl
32	277.4	81.8	405	17	US-10-309-764-62	Sequence 62, Appl
33	276.8	81.7	414	15	US-10-325-694-143	Sequence 143, App
34	276.6	81.6	369	21	US-10-625-307A-7	Sequence 7, Appli
35	276.2	81.5	397	9	US-09-878-178-2106	Sequence 2106, Ap
36	276.2	81.5	397	13	US-10-046-935-2106	Sequence 2106, Ap
37	276.2	81.5	397	14	US-10-146-502-2106	Sequence 2106, Ap
38	275.8	81.4	405	17	US-10-309-764-78	Sequence 78, Appl
39	275.8	81.4	405	17	US-10-309-764-138	Sequence 138, App
40	275	81.1	345	21	US-10-625-307A-5	Sequence 5, Appli
41	275	81.1	345	21	US-10-625-307A-60	Sequence 60, Appl
42	275	81.1	350	21	US-10-625-307A-36	Sequence 36, Appl
43	274.6	81.0	354	15	US-10-324-493-7	Sequence 7, Appli
44	274.6	81.0	366	21	US-10-727-155-33	Sequence 33, Appl
45	274.4	80.9	376	17	US-10-292-088-73	Sequence 73, Appl

ALIGNMENTS

RESULT 1
US-10-325-694-145
; Sequence 145, Application US/10325694
; Publication No. US20030148463A1
; GENERAL INFORMATION:
; APPLICANT: KUFER, PETER
; APPLICANT: RAUM, TOBIAS
; TITLE OF INVENTION: NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN
; TITLE OF INVENTION: RECEPTORS AND USES THEREOF
; FILE REFERENCE: 38164000
; CURRENT APPLICATION NUMBER: US/10/325,694
; CURRENT FILING DATE: 2002-12-19
; PRIOR APPLICATION NUMBER: US/09/403,107
; PRIOR FILING DATE: 1999-10-14
; NUMBER OF SEQ ID NOS: 152
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 145
; LENGTH: 372
; TYPE: DNA
; ORGANISM: HUMAN
US-10-325-694-145

Query Match 100.0%; Score 339; DB 15; Length 372;
Best Local Similarity 100.0%; Pred. No. 2e-102;
Matches 339; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60

```
Qy          61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120

Qy         121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180

Qy         181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240

Qy         241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300

Qy         301 GGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
               |||||||||||||||||||||||||||||||||||||||
Db         301 GGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339


RESULT 2
US-10-406-830-20
; Sequence 20, Application US/10406830
; Publication No. US20040071696A1
; GENERAL INFORMATION:
;  APPLICANT: ADAMS, GREGORY P.
;  APPLICANT:  HORAK, EVA M.
;  APPLICANT:  WEINER, LOUIS M.
;  APPLICANT:  JAMES, MARKS D.
;  TITLE OF INVENTION: BISPECIFIC SINGLE CHAIN Fv ANTIBODY MOLECULES AND METHODS OF USE
;  TITLE OF INVENTION:  THEREOF
;  FILE REFERENCE: 407T-000410US
;  CURRENT APPLICATION NUMBER: US/10/406,830
;  CURRENT FILING DATE:  2003-04-04
;  PRIOR APPLICATION NUMBER: US 60/370,276
;  PRIOR FILING DATE: 2002-04-05
;  NUMBER OF SEQ ID NOS: 37
;  SOFTWARE: PatentIn version 3.2
; SEQ ID NO 20
;   LENGTH: 729
;   TYPE: DNA
;   ORGANISM: Artificial
;   FEATURE:
;   OTHER INFORMATION: Nucleic acid encoding single chain antibody.
US-10-406-830-20

  Query Match             88.8%;  Score 301.2;  DB 18;  Length 729;
  Best Local Similarity   93.2%;  Pred. No. 1e-89;
  Matches  315;  Conservative    0;  Mismatches   23;  Indels    0;  Gaps    0;

Qy           2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
               |||||||||| | ||||||||||||| |||||| ||||||||| ||||||||||||||||
Db           8 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 67

Qy          62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
               |||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||||
Db          68 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 127

Qy         122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
               |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         128 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 187

Qy         182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
               |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         188 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 247

Qy         242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
               |||||||||||||||||||||||||||||||||||| ||||||||||||||   |
Db         248 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGAGTATTACTGTGCGAAGTATCCTT 307

Qy         302 GCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                  |||||||||||||||||||||||||||||||||||
Db         308 TAAACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 345


RESULT 3
US-10-309-764-54
; Sequence 54, Application US/10309764
; Publication No. US20030232009A1
; GENERAL INFORMATION:
;  APPLICANT: Foltz, Ian
;  APPLICANT:  Babcook, John
;  APPLICANT:  Palathumpat, Raju
;  APPLICANT:  Yang, Xiao-dong
;  APPLICANT:  King, Chadwick T.
;  TITLE OF INVENTION: ANTI-CDR45RB ANTIBODIES FOR USE IN
;  TITLE OF INVENTION:  TREATING AUTOIMMUNE DISEASE AND TRANSPLANT REJECTION
;  FILE REFERENCE: ABGENIX.029A
;  CURRENT APPLICATION NUMBER: US/10/309,764
;  CURRENT FILING DATE:  2002-12-02
;  PRIOR APPLICATION NUMBER: 60/337,276
;  PRIOR FILING DATE: 2001-12-03
;  NUMBER OF SEQ ID NOS: 147
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 54
;   LENGTH: 396
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-309-764-54

  Query Match             87.0%;  Score 294.8;  DB 17;  Length 396;
  Best Local Similarity   92.0%;  Pred. No. 1.1e-87;
  Matches  311;  Conservative    0;  Mismatches   27;  Indels    0;  Gaps    0;

Qy           2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
               |||||||||| | ||||||||||||| |||||| ||||||||| ||||||||||||||||
Db          59 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 118

Qy          62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
               |||||||||| ||||||||||||||    | | |||| |||||||||||||||||||||||||
Db         119 CCTGTGCAGCGTCTGGATTCACCTTCAGTAACTATGGCATGCACTGGGTCCGCCAGGCTC 178

Qy         122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
               |||||||||||||||||||||||| |||||||  |||||||||||||| | || |||||
Db         179 CAGGCAAGGGGCTGGAGTGGGTGGCTGTTATATGGTATGATGGAAGTAAGAGATTCTATG 238

Qy         182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
               ||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         239 CAGGCTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 298

Qy         242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
               |||||||||||||||||||||||||||||||||||| ||||||||||||||| ||    ||
Db         299 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAACTACAG 358

Qy         302 GCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                 |||||||||||||||||| ||||||||||||||||
Db         359 ACTACTGGGGCCAGGGAACCATGGTCACCGTCTCCTCA 396


RESULT 4
US-10-738-120-9
; Sequence 9, Application US/10738120
; Publication No. US20040208873A1
; GENERAL INFORMATION:
;  APPLICANT: Teeling, Jessica
;  APPLICANT:  Parren, Paul
;  APPLICANT:  Baadsgaard, Ole
;  APPLICANT:  Hudson, Debra
;  APPLICANT:  Petersen, Jorgen
;  TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES AGAINST INTERLEUKIN 8 (IL-8)
;  FILE REFERENCE: MXI-278
;  CURRENT APPLICATION NUMBER: US/10/738,120
;  CURRENT FILING DATE:  2003-12-16
;  PRIOR APPLICATION NUMBER: 60/433,728
;  PRIOR FILING DATE: 2002-12-16
```

```
;  NUMBER OF SEQ ID NOS: 45
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 9
;   LENGTH: 342
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-738-120-9

  Query Match             85.3%;  Score 289.2;  DB 20;  Length 342;
  Best Local Similarity   92.6%;  Pred. No. 7.8e-86;
  Matches  315;  Conservative    0;  Mismatches   23;  Indels    2;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||| |||||| |||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| |||||||||||||   |   |||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCGTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||  |||||||||||||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATGGTATGATGGAAGTAATAAATACTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGC--GAAAAAGGA 299
              |||||||||||||||||||| |||||||||||||||||||||||||  || | | |
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGATACTT 301

Qy        300 AGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
              | ||||||||||||||||||||||||||||||||||||||
Db        302 TGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 341


RESULT 5
US-10-805-177-25
; Sequence 25, Application US/10805177
; Publication No. US20050084449A1
; GENERAL INFORMATION:
;  APPLICANT: Landes, Gregory M.
;  APPLICANT:  Chen, Francine
;  APPLICANT:  Bezabeh, Binyam
;  APPLICANT:  Foltz, Ian
;  APPLICANT:  Tse, Kam Fai
;  APPLICANT:  Jeffers, Michael
;  APPLICANT:  Mesri, Mehdi
;  APPLICANT:  Starling, Gary
;  APPLICANT:  Mezes, Peter
;  APPLICANT:  Khramtsov, Nikolia
;  TITLE OF INVENTION: ANTIBODIES AGAINST T CELL IMMUNOGLOBULIN
;  TITLE OF INVENTION:  DOMAIN AND MUCIN DOMAIN 1 (TIM-1) ANTIGEN AND USES THEREOF
;  FILE REFERENCE: ABXCUR.006A
;  CURRENT APPLICATION NUMBER: US/10/805,177
;  CURRENT FILING DATE:  2004-03-19
;  PRIOR APPLICATION NUMBER: 60/456,652
;  PRIOR FILING DATE: 2003-03-19
;  NUMBER OF SEQ ID NOS: 141
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 25
;   LENGTH: 537
;   TYPE: DNA
;   ORGANISM: Homo Sapiens
US-10-805-177-25

  Query Match             84.1%;  Score 285.2;  DB 21;  Length 537;
  Best Local Similarity   90.2%;  Pred. No. 2e-84;
  Matches  305;  Conservative    0;  Mismatches   33;  Indels    0;  Gaps    0;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              ||||||||||||    |||| ||||||| |||||| |||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGAGCAGTCGGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| |||||||||||| |||| |||||||||||||||||||||||||||||||
Db         62 CCTGTGCAGCGTCTGGATTCACCTTCACTAACTATGGCTTGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||| |||||||||||||  |||||||||| |||||| ||||||||||
Db        122 CAGGCAAGGGGCTGGATTGGGTGGCAGTTATATGGTATGATGGAAGTCATAAATTCTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||| | ||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTCTTTC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              |||||||||||||||||||||| |||||||||||||||||||||||| || | |    |
Db        242 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTACGCGAGATCTTG 301

Qy        302 GCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
              |||||||||||||||||||||||||||||||||||||||
Db        302 ACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339


RESULT 6
US-10-010-729-8
; Sequence 8, Application US/10010729
; Publication No. US20030185827A1
; GENERAL INFORMATION:
;  APPLICANT: Rodriguez, Moses
;  APPLICANT:  Miller, David J.
;  APPLICANT:  Pease, Larry R.
;  TITLE OF INVENTION: Human IgM Antibodies and Diagnostic and
;  TITLE OF INVENTION:  Therapeutic Uses Thereof Particularly in the Central Nervous
;  TITLE OF INVENTION:  System
;  FILE REFERENCE: 1199-1-005CIP2
;  CURRENT APPLICATION NUMBER: US/10/010,729
;  CURRENT FILING DATE:  2001-11-13
;  PRIOR APPLICATION NUMBER: 09/730,473
;  PRIOR FILING DATE: 2000-12-05
;  PRIOR APPLICATION NUMBER: 09/580,787
;  PRIOR FILING DATE: 2000-05-30
;  PRIOR APPLICATION NUMBER: 09/322,862
;  PRIOR FILING DATE: 1999-05-28
;  PRIOR APPLICATION NUMBER: 08/779,784
;  PRIOR FILING DATE: 1997-01-07
;  PRIOR APPLICATION NUMBER: 08/692,084
;  PRIOR FILING DATE: 1996-08-08
;  PRIOR APPLICATION NUMBER: 08/236,520
;  PRIOR FILING DATE: 1994-04-29
;  NUMBER OF SEQ ID NOS: 80
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 8
;   LENGTH: 357
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-010-729-8

  Query Match             83.9%;  Score 284.4;  DB 16;  Length 357;
  Best Local Similarity   90.4%;  Pred. No. 3.2e-84;
  Matches  322;  Conservative    0;  Mismatches   16;  Indels   18;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||| | |||||||||||| |||||| |||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |   |||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
```

```
Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGG--- 298
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||| |||
Db        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAGAGGTGA 301

Qy        299 ---------------AAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                            | ||||||||||||||||||||||||||||||||||||||
Db        302 CTGCTATTCCCTACTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 357


RESULT 7
US-10-638-265-17
; Sequence 17, Application US/10638265
; Publication No. US20050031614A1
; GENERAL INFORMATION:
;  APPLICANT: Roskos, Lorin
;  APPLICANT:  Foltz, Ian
;  APPLICANT:  King, Chadwick
;  TITLE OF INVENTION: ANTIBODIES DIRECTED TO PARATHYROID
;  TITLE OF INVENTION:  HORMONE (PTH) AND USES THEREOF
;  FILE REFERENCE: ABGENIX.092A
;  CURRENT APPLICATION NUMBER: US/10/638,265
;  CURRENT FILING DATE:  2003-08-08
;  NUMBER OF SEQ ID NOS: 97
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 17
;   LENGTH: 352
;   TYPE: DNA
;   ORGANISM: Homosapien
US-10-638-265-17

  Query Match             83.8%;  Score 284;  DB 21;  Length 352;
  Best Local Similarity   90.9%;  Pred. No. 4.3e-84;
  Matches  318;  Conservative    0;  Mismatches   20;  Indels   12;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||||||| | |||||||||||||| |||||| |||||||| ||||||||||||||||
Db          2 AGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 61

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||||
Db         62 CCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCTC 121

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||| |||||||||||||||||||||||||||||||||||| |||||
Db        122 CAGGCAAGGGGCTGGACTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATTCTATG 181

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241

Db        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGA-- 299
Qy        300 ||||||||||||||||||||||||||||||||||||||||||||||||||||||| | | |
              AATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAGAGACCATT 301
Db        302 GGGAGCTAAGGCTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
                      ||||||||||||||||||||||||||||||||||||||
                      ACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 351

RESULT 8
US-10-309-764-58
; Sequence 58, Application US/10309764
; Publication No. US20030232009A1
; GENERAL INFORMATION:
;  APPLICANT: Foltz, Ian
;  APPLICANT:  Babcook, John
;  APPLICANT:  Palathumpat, Raju
;  APPLICANT:  Yang, Xiao-dong
;  APPLICANT:  King, Chadwick T.
;  TITLE OF INVENTION: ANTI-CDR45RB ANTIBODIES FOR USE IN
;  TITLE OF INVENTION:  TREATING AUTOIMMUNE DISEASE AND TRANSPLANT REJECTION
;  FILE REFERENCE: ABGENIX.029A
;  CURRENT APPLICATION NUMBER: US/10/309,764
;  CURRENT FILING DATE:  2002-12-02
;  PRIOR APPLICATION NUMBER: 60/337,276
;  PRIOR FILING DATE: 2001-12-03
;  NUMBER OF SEQ ID NOS: 147
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 58
;   LENGTH: 405
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-309-764-58

  Query Match             83.7%;  Score 283.8;  DB 17;  Length 405;
  Best Local Similarity   91.1%;  Pred. No. 5.2e-84;
  Matches  316;  Conservative    0;  Mismatches   22;  Indels    9;  Gaps    1;

Qy          2 AGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTCT 61
              |||||||| || | |||||||||||||| |||||| |||||||| ||||||||||||||||
Db         59 AGGTGCAGTTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCT 118

Qy         62 CCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCTC 121
              |||||||||| ||||||||||||||    | | |||| ||||||||||||||||||||||||
Db        119 CCTGTGCAGCGTCTGGATTCACCTTCAGTAACTATGGCATGCACTGGGTCCGCCAGGCTC 178

Qy        122 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTATG 181
              |||||||||||||||||||||||||||||||||||  ||||||||||||||||||||||||
Db        179 CAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATGGTATGATGGAAGTAATAAATACTATG 238

Qy        182 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATC 241
              |||||||||||||||||||||||||||||||||||||||||||||||||||| |||||||||
Db        239 CAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCGAAGAACACGCTGTATC 298

Qy        242 TGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAAG 301
              |||||||||||||||||||||| |||||||||||||||||||||||||||| |  ||  |
Db        299 TGCAAATGAACAGCCTGAGAGCCGAGGACACGGCTGTGTATTACTGTGCGAGAGGGGGGG 358

Qy        302 G---------CTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 339
              |         ||||||||||||||||||||||||||||||||||||||
Db        359 GTGACTTTGACTACTGGGGCCAGGGAACCCTGGTCACCGTCTCCTCA 405


RESULT 9
US-10-642-120-5
; Sequence 5, Application US/10642120
; Publication No. US20040131610A1
; GENERAL INFORMATION:
;  APPLICANT: Thorpe, Philip E.
;  APPLICANT:  Soares, M. Melina
;  APPLICANT:  Ran, Sophia
;  TITLE OF INVENTION: Methods for Treating Viral Infections Using Antibodies to
;  TITLE OF INVENTION:  Aminophospholipids
;  FILE REFERENCE: 4001.002900
;  CURRENT APPLICATION NUMBER: US/10/642,120
;  CURRENT FILING DATE:  2003-08-15
;  PRIOR APPLICATION NUMBER: US 10/621,269
;  PRIOR FILING DATE: 2003-07-15
;  PRIOR APPLICATION NUMBER: 60/396,263
;  PRIOR FILING DATE: 2002-07-15
;  NUMBER OF SEQ ID NOS: 9
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
;   LENGTH: 783
;   TYPE: DNA
;   ORGANISM: ARTIFICIAL SEQUENCE
;   FEATURE:
;   OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-10-642-120-5

  Query Match             83.2%;  Score 282;  DB 19;  Length 783;
  Best Local Similarity   91.7%;  Pred. No. 2.6e-83;
```

```
  Matches  310;  Conservative    0;  Mismatches   25;  Indels    3;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              ||||||||||||  | ||||||||||||| |||||| ||||||||| ||||||||||||||
Db         19 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 78

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              ||||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         79 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 138

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        139 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 198

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        199 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 258

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||| | |  | |
Db        259 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCCGTGTATTACTGTGCAAGATTGCAT 318

Qy        301 GGCTA---CTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
              |   |    |||||||| || ||||||||||||||||
Db        319 GCTCAGACTTGGGGCCAAGGTACCCTGGTCACCGTCTC 356


RESULT 10
US-10-642-060-5
; Sequence 5, Application US/10642060
; Publication No. US20040131621A1
; GENERAL INFORMATION:
;  APPLICANT: Thorpe, Philip E.
;  APPLICANT:  Soares, M. Melina
;  APPLICANT:  Ran, Sophia
;  TITLE OF INVENTION: Combinations and Kits for Treating Viral Infections Using Antibod
;  TITLE OF INVENTION:  Aminophospholipids
;  FILE REFERENCE: 4001.002982
;  CURRENT APPLICATION NUMBER: US/10/642,060
;  CURRENT FILING DATE:  2003-08-15
;  PRIOR APPLICATION NUMBER: US 10/621,269
;  PRIOR FILING DATE: 2003-07-15
;  PRIOR APPLICATION NUMBER: 60/396,263
;  PRIOR FILING DATE: 2002-07-15
;  NUMBER OF SEQ ID NOS: 9
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
;   LENGTH: 783
;   TYPE: DNA
;   ORGANISM: ARTIFICIAL SEQUENCE
;   FEATURE:
;   OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-10-642-060-5

  Query Match           83.2%;  Score 282;  DB 19;  Length 783;
  Best Local Similarity   91.7%;  Pred. No. 2.6e-83;
  Matches  310;  Conservative    0;  Mismatches   25;  Indels    3;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              ||||||||||||  | ||||||||||||| |||||| ||||||||| ||||||||||||||
Db         19 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 78

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              ||||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         79 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 138

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        139 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 198

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        199 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 258

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
              |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||| | |  | |
Db        259 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCCGTGTATTACTGTGCAAGATTGCAT 318

Qy        301 GGCTA---CTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
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Db        319 GCTCAGACTTGGGGCCAAGGTACCCTGGTCACCGTCTC 356


RESULT 11
US-10-642-122-5
; Sequence 5, Application US/10642122
; Publication No. US20040131622A1
; GENERAL INFORMATION:
;  APPLICANT: Thorpe, Philip E.
;  APPLICANT:  Soares, M. Melina
;  APPLICANT:  Ran, Sophia
;  TITLE OF INVENTION: Combinations and Kits for Treating Viral Infections Using
;  TITLE OF INVENTION:  Immunoconjugates to Aminophospholipids
;  FILE REFERENCE: 3999.002985
;  CURRENT APPLICATION NUMBER: US/10/642,122
;  CURRENT FILING DATE:  2003-08-15
;  PRIOR APPLICATION NUMBER: US 10/621,269
;  PRIOR FILING DATE: 2003-07-15
;  PRIOR APPLICATION NUMBER: 60/396,263
;  PRIOR FILING DATE: 2002-07-15
;  NUMBER OF SEQ ID NOS: 9
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
;   LENGTH: 783
;   TYPE: DNA
;   ORGANISM: ARTIFICIAL SEQUENCE
;   FEATURE:
;   OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-10-642-122-5

  Query Match           83.2%;  Score 282;  DB 19;  Length 783;
  Best Local Similarity   91.7%;  Pred. No. 2.6e-83;
  Matches  310;  Conservative    0;  Mismatches   25;  Indels    3;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              ||||||||||||  | ||||||||||||| |||||| ||||||||| ||||||||||||||
Db         19 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 78

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              ||||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         79 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 138

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
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Db        139 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 198

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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Db        199 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 258

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
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Db        259 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCCGTGTATTACTGTGCAAGATTGCAT 318

Qy        301 GGCTA---CTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
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Db        319 GCTCAGACTTGGGGCCAAGGTACCCTGGTCACCGTCTC 356


RESULT 12
US-10-642-124-5
; Sequence 5, Application US/10642124
; Publication No. US20040161429A1
; GENERAL INFORMATION:
;  APPLICANT: Thorpe, Philip E.
;  APPLICANT:  Soares, M. Melina
```

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;  APPLICANT:  Ran, Sophia
;  TITLE OF INVENTION: Compositions for Treating Viral Infections Using Immunoconjugates
;  TITLE OF INVENTION:  Aminophospholipids
;  FILE REFERENCE: 3999.002984
;  CURRENT APPLICATION NUMBER: US/10/642,124
;  CURRENT FILING DATE:  2003-08-15
;  PRIOR APPLICATION NUMBER: US 10/621,269
;  PRIOR FILING DATE: 2003-07-15
;  PRIOR APPLICATION NUMBER: 60/396,263
;  PRIOR FILING DATE: 2002-07-15
;  NUMBER OF SEQ ID NOS: 9
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
;   LENGTH: 783
;   TYPE: DNA
;   ORGANISM: ARTIFICIAL SEQUENCE
;   FEATURE:
;   OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-10-642-124-5

  Query Match             83.2%;  Score 282;  DB 19;  Length 783;
  Best Local Similarity   91.7%;  Pred. No. 2.6e-83;
  Matches  310;  Conservative    0;  Mismatches   25;  Indels    3;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              ||||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||
Db         19 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 78

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              ||||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         79 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 138

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        139 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 198

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        199 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 258

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
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Db        259 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCCGTGTATTACTGTGCAAGATTGCAT 318

Qy        301 GGCTA---CTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
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Db        319 GCTCAGACTTGGGGCCAAGGTACCCTGGTCACCGTCTC 356


RESULT 13
US-10-621-269-5
; Sequence 5, Application US/10621269
; Publication No. US20040170620A1
; GENERAL INFORMATION:
;  APPLICANT: Thorpe, Philip E.
;  APPLICANT:  Ran, Sophia
;  TITLE OF INVENTION: Selected Antibody Compositions for Binding to Aminophospholipids
;  FILE REFERENCE: 4001.003000
;  CURRENT APPLICATION NUMBER: US/10/621,269
;  CURRENT FILING DATE:  2003-07-15
;  PRIOR APPLICATION NUMBER: 60/396,263
;  PRIOR FILING DATE: 2002-07-15
;  NUMBER OF SEQ ID NOS: 9
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
;   LENGTH: 783
;   TYPE: DNA
;   ORGANISM: ARTIFICIAL SEQUENCE
;   FEATURE:
;   OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-10-621-269-5

  Query Match             83.2%;  Score 282;  DB 19;  Length 783;
  Best Local Similarity   91.7%;  Pred. No. 2.6e-83;
  Matches  310;  Conservative    0;  Mismatches   25;  Indels    3;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              ||||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||
Db         19 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 78

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              ||||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         79 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 138

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
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Db        139 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 198

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
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Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
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Qy        301 GGCTA---CTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
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Db        319 GCTCAGACTTGGGGCCAAGGTACCCTGGTCACCGTCTC 356


RESULT 14
US-10-620-850-5
; Sequence 5, Application US/10620850
; Publication No. US20040175378A1
; GENERAL INFORMATION:
;  APPLICANT: Thorpe, Philip E.
;  APPLICANT:  Ran, Sophia
;  TITLE OF INVENTION: Selected Antibody Compositions and Methods for Binding to
;  TITLE OF INVENTION:  Aminophospholipids
;  FILE REFERENCE: 4001.003082
;  CURRENT APPLICATION NUMBER: US/10/620,850
;  CURRENT FILING DATE:  2003-07-15
;  PRIOR APPLICATION NUMBER: 60/396,263
;  PRIOR FILING DATE: 2002-07-15
;  PRIOR APPLICATION NUMBER: 09/613,430
;  PRIOR FILING DATE: 2000-07-10
;  NUMBER OF SEQ ID NOS: 9
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
;   LENGTH: 783
;   TYPE: DNA
;   ORGANISM: ARTIFICIAL SEQUENCE
;   FEATURE:
;   OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-10-620-850-5

  Query Match             83.2%;  Score 282;  DB 19;  Length 783;
  Best Local Similarity   91.7%;  Pred. No. 2.6e-83;
  Matches  310;  Conservative    0;  Mismatches   25;  Indels    3;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              ||||||||||| | |||||||||||| |||||| ||||||||| ||||||||||||||
Db         19 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 78

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              ||||||||||||||||||||||||||    |    |||| ||||||||||||||||||||||
Db         79 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 138

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        139 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 198

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        199 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 258
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Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
              ||||||||||||||||||||||||||||||||| |||||||||||||| | |  | |
Db        259 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCCGTGTATTACTGTGCAAGATTGCAT 318

Qy        301 GGCTA---CTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
              |   |    |||||||| || ||||||||||||||||||
Db        319 GCTCAGACTTGGGGCCAAGGTACCCTGGTCACCGTCTC 356


RESULT 15
US-10-642-118-5
; Sequence 5, Application US/10642118
; Publication No. US20040208868A1
; GENERAL INFORMATION:
;  APPLICANT: Thorpe, Philip E.
;  APPLICANT:  Ran, Sophia
;  TITLE OF INVENTION: Selected Antibody CDRs for Binding to Aminophospholipids
;  FILE REFERENCE: 4001.003085
;  CURRENT APPLICATION NUMBER: US/10/642,118
;  CURRENT FILING DATE:  2003-08-15
;  PRIOR APPLICATION NUMBER: US 10/621,269
;  PRIOR FILING DATE: 2003-07-15
;  PRIOR APPLICATION NUMBER: 60/396,263
;  PRIOR FILING DATE: 2002-07-15
;  NUMBER OF SEQ ID NOS: 9
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
;   LENGTH: 783
;   TYPE: DNA
;   ORGANISM: ARTIFICIAL SEQUENCE
;   FEATURE:
;   OTHER INFORMATION: SYNTHETIC OLIGONUCLEOTIDE
US-10-642-118-5

  Query Match             83.2%;  Score 282;  DB 20;  Length 783;
  Best Local Similarity   91.7%;  Pred. No. 2.6e-83;
  Matches  310;  Conservative    0;  Mismatches   25;  Indels    3;  Gaps    1;

Qy          1 GAGGTGCAGCTGCTCGAGTCTGGGGGAGTCGTGGTACAGCCTGGGGGGTCCCTGAGACTC 60
              |||||||||||| | |||||||||||||| |||||| ||||||||| ||||||||||||||
Db         19 GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTC 78

Qy         61 TCCTGTGCAGCCTCTGGATTCACCTTTGATGATTATGCCATGCACTGGGTCCGCCAGGCT 120
              |||||||||||||||||||||||||    |   |||| ||||||||||||||||||||||
Db         79 TCCTGTGCAGCCTCTGGATTCACCTTCAGTAGCTATGGCATGCACTGGGTCCGCCAGGCT 138

Qy        121 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        139 CCAGGCAAGGGGCTGGAGTGGGTGGCAGTTATATCATATGATGGAAGTAATAAATACTAT 198

Qy        181 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        199 GCAGACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTAT 258

Qy        241 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCTGTGTATTACTGTGCGAAAAAGGAA 300
              ||||||||||||||||||||||||||||||||| |||||||||||||| | |  | |
Db        259 CTGCAAATGAACAGCCTGAGAGCTGAGGACACGGCCGTGTATTACTGTGCAAGATTGCAT 318

Qy        301 GGCTA---CTGGGGCCAGGGAACCCTGGTCACCGTCTC 335
              |   |    |||||||| || ||||||||||||||||||
Db        319 GCTCAGACTTGGGGCCAAGGTACCCTGGTCACCGTCTC 356


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Job time : 539.75 secs
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GenCore version 5.1.6

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Total number of hits satisfying chosen parameters: 9416466

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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5: gb_ov:*
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8: gb_pl:*
9: gb_pr:*
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11: gb_sts:*
12: gb_sy:*
13: gb_un:*
14: gb_vi:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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5	321	100.0	1630	6	AX023365	AX023365 Sequence
6	321	100.0	1630	6	AX023367	AX023367 Sequence
7	260.2	81.1	321	6	A84374	A84374 Sequence 14
8	260.2	81.1	321	6	BD139668	BD139668 A novel m
9	260.2	81.1	321	6	AX003761	AX003761 Sequence
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31	234.4	73.0	936	6	BD248703	BD248703 Immunoglo
32	233	72.6	324	9	HSVJTEL9	X61642 Human Ig re
33	233	72.6	450	9	AF103775	AF103775 Homo sapi
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36	231.4	72.1	324	9	AB063929	AB063929 Homo sapi
37	231.4	72.1	333	6	BD097236	BD097236 A therape
38	231.4	72.1	346	9	HSG1B6IGK	X94431 H.sapiens r
39	231.4	72.1	388	9	HSA548508	AJ548508 Homo sapi
40	231.4	72.1	720	6	CQ761230	CQ761230 Sequence
41	231.4	72.1	720	6	CQ840583	CQ840583 Sequence
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43	231.4	72.1	720	6	AX740176	AX740176 Sequence
44	231.4	72.1	720	6	BD077368	BD077368 Method of
45	231.4	72.1	761	9	AB064051	AB064051 Homo sapi

ALIGNMENTS

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RESULT 1
A84380
LOCUS       A84380                   321 bp    DNA     linear   PAT 21-JAN-2000
DEFINITION  Sequence 147 from Patent WO9846645.
ACCESSION   A84380
VERSION     A84380.1  GI:6733303
KEYWORDS    .
SOURCE      unidentified
  ORGANISM  unidentified
            unclassified.
REFERENCE   1  (bases 1 to 321)
  AUTHORS   Kufer,P. and Raum,T.
  TITLE     NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN RECEPTORS AND
            USES THEREOF
  JOURNAL   Patent: WO 9846645-A 147 22-OCT-1998;
            KUFER PETER (DE); RAUM TOBIAS (DE)
FEATURES             Location/Qualifiers
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                     /db_xref="taxon:32644"
     CDS             <1. .>321
                     /note="unnamed protein product"
                     /codon_start=1
                     /protein_id="CAB69290.1"
                     /db_xref="GI:6733304"
                     /translation="ELQMTQSPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGQPP
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                     RLDIQ"
ORIGIN

  Query Match          100.0%;  Score 321;  DB 6;  Length 321;
  Best Local Similarity  100.0%;  Pred. No. 2.3e-97;
  Matches  321;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
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Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Db        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
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Db        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
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Db        301 GGGACACGACTGGACATTCAA 321


RESULT 2
BD075296
LOCUS       BD075296                321 bp    DNA     linear   PAT 27-AUG-2002
DEFINITION  Novel method for the production of anti-human antigen receptors and
            uses thereof.
ACCESSION   BD075296
VERSION     BD075296.1  GI:22620899
KEYWORDS    JP 2001519824-A/25.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 321)
  AUTHORS   Kufer,P. and Raum,T.
  TITLE     Novel method for the production of anti-human antigen receptors and
            uses thereof
  JOURNAL   Patent: JP 2001519824-A 25 23-OCT-2001;
            MICROMET AG
COMMENT     OS   Homo sapiens (human)
            PN   JP 2001519824-A/25
            PD   23-OCT-2001
            PF   14-APR-1998 JP 1998543494
            PR   14-APR-1997 EP   97106109.8
            PI   PETER KUFER,TOBIAS RAUM
            PC   C07K16/00,C07K16/30,A61K39/395
            CC   Novel method for the production of anti-human antigen  CC
                 receptors and uses
            CC   thereof
            FH   Key             Location/Qualifiers
            FT   CDS             (1). .(321).
FEATURES              Location/Qualifiers
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                      /organism="Homo sapiens"
                      /mol_type="genomic DNA"
                      /db_xref="taxon:9606"
ORIGIN

  Query Match            100.0%;  Score 321;  DB 6;  Length 321;
  Best Local Similarity  100.0%;  Pred. No. 2.3e-97;
  Matches  321;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
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Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
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Db        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
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Db        301 GGGACACGACTGGACATTCAA 321


RESULT 3
BD222938
LOCUS       BD222938                1630 bp    DNA     linear   PAT 17-JUL-2003
DEFINITION  Heterominibodies.
ACCESSION   BD222938
VERSION     BD222938.1  GI:33032708
KEYWORDS    JP 2002521053-A/32.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1630)
  AUTHORS   Kufer,P., Dreier,T., Baeuerle,P.A., Borschert,K. and Zettl,F.
  TITLE     Heterominibodies
  JOURNAL   Patent: JP 2002521053-A 32 16-JUL-2002;
            MICROMET AG
COMMENT     OS   Homo sapiens (human)
            OS   Mus musculus (mouse)
            PN   JP 2002521053-A/32
            PD   16-JUL-2002
            PF   28-JUL-1999 JP 2000562401
            PR   28-JUL-1998 EP   98114082.5
            PI   PETER KUFER,TORSTEN DREIER,PATRICK A BAEUERLE,KATRIN
            BORSCHERT,
            PI   FLORIAN ZETTL
            PC   C12N15/09,A61K35/76,A61K38/00,A61K38/21,A61P35/00,A61P35/02,
            PC   C07K19/00,
            PC   C12N5/10,C12P21/02,G01N33/53,G01N33/53//(C12N5/10,C12R1:91),
            PC   (C12P21/02,C12R1:91),C12N15/00,C12N5/00,A61K37/02,A61K37/66,
            PC   (C12N5/00,C12R1:91)
            CC   Heterominibodies
            FH   Key             Location/Qualifiers
            FT   CDS             (39). .(1607).
FEATURES              Location/Qualifiers
     source           1. .1630
                      /organism="Homo sapiens"
                      /mol_type="genomic DNA"
                      /db_xref="taxon:9606"
ORIGIN

  Query Match            100.0%;  Score 321;  DB 6;  Length 1630;
  Best Local Similarity  100.0%;  Pred. No. 2.5e-97;
  Matches  321;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Db         96 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 155

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Db        156 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 215

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        216 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 275

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Db        276 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 335

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
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Db        336 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 395

Qy        301 GGGACACGACTGGACATTCAA 321
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Db        396 GGGACACGACTGGACATTCAA 416


RESULT 4
BD222939
LOCUS       BD222939                1630 bp    DNA     linear   PAT 17-JUL-2003
DEFINITION  Heterominibodies.
ACCESSION   BD222939
VERSION     BD222939.1  GI:33032709
KEYWORDS    JP 2002521053-A/33.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 1630)
  AUTHORS   Kufer,P., Dreier,T., Baeuerle,P.A., Borschert,K. and Zettl,F.
  TITLE     Heterominibodies
  JOURNAL   Patent: JP 2002521053-A 33 16-JUL-2002;
            MICROMET AG
COMMENT     OS   Homo sapiens (human)
            OS   Mus musculus (mouse)
            PN   JP 2002521053-A/33
            PD   16-JUL-2002
            PF   28-JUL-1999 JP 2000562401
            PR   28-JUL-1998 EP   98114082.5
            PI   PETER KUFER,TORSTEN DREIER,PATRICK A BAEUERLE,KATRIN
            BORSCHERT,
            PI   FLORIAN ZETTL
            PC   C12N15/09,A61K35/76,A61K38/00,A61K38/21,A61P35/00,A61P35/02,
            PC   C07K19/00,
            PC   C12N5/10,C12P21/02,G01N33/53,G01N33/53//(C12N5/10,C12R1:91),
            PC   (C12P21/02,C12R1:91),C12N15/00,C12N5/00,A61K37/02,A61K37/66,
            PC   (C12N5/00,C12R1:91)
            CC   Heterominibodies
            FH   Key             Location/Qualifiers
            FT   CDS             (39). .(1610).
FEATURES              Location/Qualifiers
     source           1. .1630
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                      /mol_type="genomic DNA"
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ORIGIN

  Query Match             100.0%;  Score 321;  DB 6;  Length 1630;
  Best Local Similarity   100.0%;  Pred. No. 2.5e-97;
  Matches  321;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Db         96 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 155

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Db        156 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 215

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        216 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 275

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Db        276 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 335

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
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Db        336 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 395

Qy        301 GGGACACGACTGGACATTCAA 321
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Db        396 GGGACACGACTGGACATTCAA 416


RESULT 5
AX023365
LOCUS       AX023365                1630 bp    DNA     linear   PAT 15-SEP-2000
DEFINITION  Sequence 36 from Patent WO0006605.
ACCESSION   AX023365
VERSION     AX023365.1  GI:10183777
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Kufer,P., Zettl,F., Dreier,T., Baeuerle,P.A. and Borschert,K.
  TITLE     Heterominibodies
  JOURNAL   Patent: WO 0006605-A 36 10-FEB-2000;
            KUFER PETER (DE) ; ZETTL FLORIAN (DE) ; DREIER TORSTEN (DE) ;
            BAEUERLE PATRICK A (DE) ; BORSCHERT KATRIN (DE) ; MICROMET GES FUER
            BIOMEDIZINIS (DE)
FEATURES              Location/Qualifiers
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                      LQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTSGGGGSAP
                      ARSPSPSTQPWEHVNAIQEARRLLNLSRDTAAEMNETVEVISEMFDLQEPTCLQTRLE
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ORIGIN

  Query Match             100.0%;  Score 321;  DB 6;  Length 1630;
  Best Local Similarity   100.0%;  Pred. No. 2.5e-97;
  Matches  321;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Db         96 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 155

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Db        156 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 215

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        216 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 275

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Db        276 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 335

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
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Qy        301 GGGACACGACTGGACATTCAA 321
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Db        396 GGGACACGACTGGACATTCAA 416


RESULT 6
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AX023367
LOCUS       AX023367               1630 bp    DNA     linear   PAT 15-SEP-2000
DEFINITION  Sequence 38 from Patent WO0006605.
ACCESSION   AX023367
VERSION     AX023367.1  GI:10183779
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Kufer,P., Zettl,F., Dreier,T., Baeuerle,P.A. and Borschert,K.
  TITLE     Heterominibodies
  JOURNAL   Patent: WO 0006605-A 38 10-FEB-2000;
            KUFER PETER (DE) ; ZETTL FLORIAN (DE) ; DREIER TORSTEN (DE) ;
            BAEUERLE PATRICK A (DE) ; BORSCHERT KATRIN (DE) ; MICROMET GES FUER
            BIOMEDIZINIS (DE)
FEATURES             Location/Qualifiers
     source          1. .1630
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     CDS             39. .1613
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                     /protein_id="CAC08836.1"
                     /db_xref="GI:10183780"
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                     HTRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESV
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                     LNRWITFCQSIISTLT"
ORIGIN

  Query Match           100.0%;  Score 321;  DB 6;  Length 1630;
  Best Local Similarity  100.0%;  Pred. No. 2.5e-97;
  Matches  321;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         96 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 155

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        156 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 215

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        216 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 275

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db        336 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 395

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||||||||
Db        396 GGGACACGACTGGACATTCAA 416


RESULT 7
A84374
LOCUS       A84374                 321 bp    DNA     linear   PAT 21-JAN-2000
DEFINITION  Sequence 141 from Patent WO9846645.
ACCESSION   A84374
VERSION     A84374.1  GI:6733297
KEYWORDS    .
SOURCE      unidentified
  ORGANISM  unidentified
            unclassified.
REFERENCE   1  (bases 1 to 321)
  AUTHORS   Kufer,P. and Raum,T.
  TITLE     NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN RECEPTORS AND
            USES THEREOF
  JOURNAL   Patent: WO 9846645-A 141 22-OCT-1998;
            KUFER PETER (DE); RAUM TOBIAS (DE)
FEATURES             Location/Qualifiers
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                     /codon_start=1
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                     /db_xref="GI:6733298"
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                     KLEIK"
ORIGIN

  Query Match           81.1%;  Score 260.2;  DB 6;  Length 321;
  Best Local Similarity  88.2%;  Pred. No. 1e-76;
  Matches  283;  Conservative    0;  Mismatches   38;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||||||||||||||||||||||||||||||||||||||||||||| ||||| ||||||||||||||||
Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCTTCTGTGGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||| ||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGTCGGACAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              |||||||| ||||| |  |||||||||| ||| ||||||||||||||||||| || || |||
Db        181 CGATTCAGTGGCAGCGGGTCTGGGACAGATTTCACTCTCACCATCAGCAGTCTACAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||| ||| |||||||  ||||| |||  | ||    | |||  ||| || ||||||
Db        241 GAAGATTCTGCAACTTACTACTGTCAGCAGAGTTACGACATCCCGTACACTTTTGGCCAG 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||    ||||| ||  ||
Db        301 GGGACCAAGCTGGAGATCAAA 321


RESULT 8
BD139668
LOCUS       BD139668               321 bp    DNA     linear   PAT 18-SEP-2002
DEFINITION  A novel method of identifying binding site domains that retain the
            capacity of binding to an epitope.
ACCESSION   BD139668
VERSION     BD139668.1  GI:23234613
KEYWORDS    JP 2002508924-A/53.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 321)
  AUTHORS   Kufer,P., Raum,T., Borschert,K., Zettl,F. and Lutterbuese,R.
  TITLE     A novel method of identifying binding site domains that retain the
            capacity of binding to an epitope
  JOURNAL   Patent: JP 2002508924-A 53 26-MAR-2002;
            PETER KUFER
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COMMENT     OS   Homo sapiens (human)
            PN   JP 2002508924-A/53
            PD   26-MAR-2002
            PF   16-NOV-1998 JP 2000521184
            PR   17-NOV-1997 EP   97120096.9
            PI   PETER KUFER,TOBIAS RAUM,KATRIN BORSCHERT,FLORIAN ZETTL,RALF
            PI   LUTTERBUESE
            PC   C12N15/09,A61K38/00,A61K38/22,A61K38/43,A61K39/395,C07K14/705,
            PC   C07K16/30,
            PC   C12N1/21,C12N5/10,C12P21/02,C12P21/08,G01N33/566,C12N15/00, PC
              A61K37/02,
            PC   A61K37/24,A61K37/48,C12N5/00
            CC   A novel method of identifying binding site
            domains that retain
            CC             the
            CC   capacity of binding to an epitope
            FH   Key             Location/Qualifiers
            FT   source          1. .321
            FT                   /organism='Homo sapiens (human)'.
FEATURES              Location/Qualifiers
     source           1. .321
                      /organism="Homo sapiens"
                      /mol_type="genomic DNA"
                      /db_xref="taxon:9606"
ORIGIN

  Query Match             81.1%;  Score 260.2;  DB 6;  Length 321;
  Best Local Similarity   88.2%;  Pred. No. 1e-76;
  Matches  283;  Conservative    0;  Mismatches   38;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||||||||||||||||||||||||||||||||||| ||||| ||||||||||||||||
Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCTTCTGTGGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||| ||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGTCGGACAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              ||||||||| ||||| |  |||||||||| ||| ||||||||||||||||| || || |||
Db        181 CGATTCAGTGGCAGCGGGTCTGGGACAGATTTCACTCTCACCATCAGCAGTCTACAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||| ||| |||||||  ||||| |||  | ||    | |||  ||| || |||||
Db        241 GAAGATTCTGCAACTTACTACTGTCAGCAGAGTTACGACATCCCGTACACTTTTGGCCAG 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||    ||||| ||  ||
Db        301 GGGACCAAGCTGGAGATCAAA 321


RESULT 9
AX003761
LOCUS       AX003761                321 bp    DNA     linear   PAT 24-AUG-2000
DEFINITION  Sequence 55 from Patent WO9925818.
ACCESSION   AX003761
VERSION     AX003761.1  GI:9927567
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1
  AUTHORS   Kufer,P. and Raum,T.
  TITLE     Method of identifying binding site domains that retain the capacity
            of binding to an epitope
  JOURNAL   Patent: WO 9925818-A 55 27-MAY-1999;
            KUFER PETER (DE); RAUM TOBIAS (DE)
FEATURES              Location/Qualifiers
     source           1. .321
                      /organism="Homo sapiens"
                      /mol_type="unassigned DNA"
                      /db_xref="taxon:9606"
ORIGIN

  Query Match             81.1%;  Score 260.2;  DB 6;  Length 321;
  Best Local Similarity   88.2%;  Pred. No. 1e-76;
  Matches  283;  Conservative    0;  Mismatches   38;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||||||||||||||||||||||||||||||||||| ||||| ||||||||||||||||
Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCTTCTGTGGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||| ||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGTCGGACAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              ||||||||| ||||| |  |||||||||| ||| ||||||||||||||||| || || |||
Db        181 CGATTCAGTGGCAGCGGGTCTGGGACAGATTTCACTCTCACCATCAGCAGTCTACAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||| ||| |||||||  ||||| |||  | ||    | |||  ||| || |||||
Db        241 GAAGATTCTGCAACTTACTACTGTCAGCAGAGTTACGACATCCCGTACACTTTTGGCCAG 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||    ||||| ||  ||
Db        301 GGGACCAAGCTGGAGATCAAA 321


RESULT 10
BD075293
LOCUS       BD075293                321 bp    DNA     linear   PAT 27-AUG-2002
DEFINITION  Novel method for the production of anti-human antigen receptors and
            uses thereof.
ACCESSION   BD075293
VERSION     BD075293.1  GI:22620896
KEYWORDS    JP 2001519824-A/22.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 321)
  AUTHORS   Kufer,P. and Raum,T.
  TITLE     Novel method for the production of anti-human antigen receptors and
            uses thereof
  JOURNAL   Patent: JP 2001519824-A 22 23-OCT-2001;
            MICROMET AG
COMMENT     OS   Homo sapiens (human)
            PN   JP 2001519824-A/22
            PD   23-OCT-2001
            PF   14-APR-1998 JP 1998543494
            PR   14-APR-1997 EP   97106109.8
            PI   PETER KUFER,TOBIAS RAUM
            PC   C07K16/00,C07K16/30,A61K39/395
            CC   Novel method for the production of anti-human antigen  CC
              receptors and uses
            CC   thereof
            FH   Key             Location/Qualifiers
            FT   CDS             (1). .(321).
FEATURES              Location/Qualifiers
     source           1. .321
                      /organism="Homo sapiens"
                      /mol_type="genomic DNA"
                      /db_xref="taxon:9606"
ORIGIN

  Query Match             81.1%;  Score 260.2;  DB 6;  Length 321;
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  Best Local Similarity   88.2%;  Pred. No. 1e-76;
  Matches  283;  Conservative    0;  Mismatches   38;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||||||||||||||||||||||||||||||||||||||| ||||| ||||||||||||||||
Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCTTCTGTGGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||| ||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGTCGGACAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              |||||||| ||||| |  |||||||||| ||| ||||||||||||||||| || || |||
Db        181 CGATTCAGTGGCAGCGGGTCTGGGACAGATTTCACTCTCACCATCAGCAGTCTACAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||| ||| ||||||||  ||||| |||  | ||    | |||  ||| || |||||
Db        241 GAAGATTCTGCAACTTACTACTGTCAGCAGAGTTACGACATCCCGTACACTTTTGGCCAG 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||    ||||| ||  ||
Db        301 GGGACCAAGCTGGAGATCAAA 321


RESULT 11
BD187178
LOCUS       BD187178                324 bp    DNA     linear   PAT 17-JUN-2003
DEFINITION  Human-typed antibody against blood coagulation factor VIII.
ACCESSION   BD187178
VERSION     BD187178.1  GI:31879467
KEYWORDS    WO 02101040-A/7.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 324)
  AUTHORS   Nakashima,T. and Yuguchi,M.
  TITLE     Human-typed antibody against blood coagulation factor VIII
  JOURNAL   Patent: WO 02101040-A 7 19-DEC-2002;
            JURIDICAL FOUNDATION THE CHEMO SERO THERAPEUTIC RESEARCH INSTITUTE,
            TOSHIHIRO NAKASHIMA,MASATO YUGUCHI
COMMENT     OS   Homo sapiens (human)
            PN   WO 02101040-A/7
            PD   19-DEC-2002
            PF   11-JUN-2002 WO 2002JP005783
            PR   12-JUN-2001 JP 01P   177640
            PI   TOSHIHIRO NAKASHIMA,MASATO YUGUCHI
            PC   C12N15/09,C07K16/14,C07K7/06,C07K7/08,C12P21/08,G01N33/53 CC
            Human-typed antibody against blood coagulation factor VIII FH   Key
                     Location/Qualifiers
            FT   source          1. .324
            FT                   /organism='Homo sapiens (human)'.
FEATURES             Location/Qualifiers
     source          1. .324
                     /organism="Homo sapiens"
                     /mol_type="genomic DNA"
                     /db_xref="taxon:9606"
ORIGIN

  Query Match             76.6%;  Score 245.8;  DB 6;  Length 324;
  Best Local Similarity   85.4%;  Pred. No. 8e-72;
  Matches  274;  Conservative    0;  Mismatches   47;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||||| || || ||    ||||| |    | ||   ||||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| ||||||| |||||||||| ||| ||||||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||||  | |||||   ||||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 12
BD187179
LOCUS       BD187179                324 bp    DNA     linear   PAT 17-JUN-2003
DEFINITION  Human-typed antibody against blood coagulation factor VIII.
ACCESSION   BD187179
VERSION     BD187179.1  GI:31879468
KEYWORDS    WO 02101040-A/8.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 324)
  AUTHORS   Nakashima,T. and Yuguchi,M.
  TITLE     Human-typed antibody against blood coagulation factor VIII
  JOURNAL   Patent: WO 02101040-A 8 19-DEC-2002;
            JURIDICAL FOUNDATION THE CHEMO SERO THERAPEUTIC RESEARCH INSTITUTE,
            TOSHIHIRO NAKASHIMA,MASATO YUGUCHI
COMMENT     OS   Homo sapiens (human)
            PN   WO 02101040-A/8
            PD   19-DEC-2002
            PF   11-JUN-2002 WO 2002JP005783
            PR   12-JUN-2001 JP 01P   177640
            PI   TOSHIHIRO NAKASHIMA,MASATO YUGUCHI
            PC   C12N15/09,C07K16/14,C07K7/06,C07K7/08,C12P21/08,G01N33/53 CC
            Human-typed antibody against blood coagulation factor VIII FH   Key
                     Location/Qualifiers
            FT   source          1. .324
            FT                   /organism='Homo sapiens (human)'.
FEATURES             Location/Qualifiers
     source          1. .324
                     /organism="Homo sapiens"
                     /mol_type="genomic DNA"
                     /db_xref="taxon:9606"
ORIGIN

  Query Match             76.6%;  Score 245.8;  DB 6;  Length 324;
  Best Local Similarity   85.4%;  Pred. No. 8e-72;
  Matches  274;  Conservative    0;  Mismatches   47;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||||| || || ||    ||||| |    | ||   ||||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| ||||||| |||||||||| ||| ||||||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240
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Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||||| |||||||  |||||||||  | |||||   ||||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 13
AB006842
LOCUS       AB006842                339 bp    mRNA    linear   PRI 09-SEP-1997
DEFINITION  Homo sapiens mRNA for HRV Fab N6-VL, partial cds.
ACCESSION   AB006842
VERSION     AB006842.1  GI:2385484
KEYWORDS    HRV Fab N6-VL.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (sites)
  AUTHORS   Itoh,K. and Suzuki,T.
  TITLE     Human anti-rotavirus Fabs
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 339)
  AUTHORS   Itoh,K.
  TITLE     Direct Submission
  JOURNAL   Submitted (26-AUG-1997) Kunihiko Itoh, Akita University Hospital,
            Pharmaceutical Science; Hondo 1-1-1, Akita, Akita 010, Japan
            (E-mail:itohk@hos.akita-u.ac.jp, Tel:+81-188-34-1111,
            Fax:+81-188-36-2628)
FEATURES             Location/Qualifiers
     source          1. .339
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /tissue_type="PBL"
     CDS             1. .>339
                     /codon_start=1
                     /product="HRV Fab N6-VL"
                     /protein_id="BAA22181.1"
                     /db_xref="GI:2385485"
                     /translation="MAELVMTQSPSSLAASVGDRVTFTCRASQSISSYLNWYQQKPGK
                     APKLLIYAASSLQSGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPITFGQ
                     GTRLEIKRTVA"
ORIGIN

  Query Match            76.1%;  Score 244.2;  DB 9;  Length 339;
  Best Local Similarity  85.0%;  Pred. No. 2.8e-71;
  Matches  273;  Conservative    0;  Mismatches   48;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||||||  |||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db          7 GAGCTCGTGATGACCCAGTCTCCATCCTCCCTGGCTGCATCTGTAGGAGACAGAGTCACC 66

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         67 TTCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 126

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||   |  | ||||||| || || ||    ||||| |    | ||   |||||||
Db        127 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 186

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| |||
Db        187 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 246

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||||| |||||||  |||||||||  | |||||   ||||||||||||||||||
Db        247 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACTCCGATCACCTTCGGCCAA 306

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||| ||| ||
Db        307 GGGACACGACTGGAGATTAAA 327


RESULT 14
BD097614
LOCUS       BD097614                342 bp    DNA     linear   PAT 27-AUG-2002
DEFINITION  Antibody library.
ACCESSION   BD097614
VERSION     BD097614.1  GI:22643188
KEYWORDS    WO 0162907-A/69.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 342)
  AUTHORS   Kurosawa,Y., Akahori,Y., Iba,Y., Morino,K., Shinohara,M.,
            Takahashi,M., Okuno,Y. and Shiraki,K.
  TITLE     Antibody library
  JOURNAL   Patent: WO 0162907-A 69 30-AUG-2001;
            MEDICAL & BIOLOGICAL LABORATORIES CO LTD,YOSHIKAZU KUROSAWA,
            YASUSHI AKAHORI,YOSHITAKA IBA,KAZUHIKO MORINO,MIDORI SHINOHARA,
            MOTOHIDE TAKAHASHI,YOSHINOBU OKUNO,KIMIYASU SHIRAKI
COMMENT     OS   Homo sapiens (human)
            PN   WO 0162907-A/69
            PD   30-AUG-2001
            PF   22-FEB-2001 WO 2001JP001298
            PR   22-FEB-2000 JP 00P    50543
            PI   YOSHIKAZU KUROSAWA,YASUSHI AKAHORI,YOSHITAKA IBA,KAZUHIKO  PI
             MORINO,
            PI   MIDORI SHINOHARA,MOTOHIDE TAKAHASHI,YOSHINOBU OKUNO,KIMIYASU
            PI   SHIRAKI
            PC   C12N15/09,C07K16/00//C12P21/08
            CC   Antibody library
            FH   Key             Location/Qualifiers
            FT   source          1. .342
            FT                   /organism='Homo sapiens (human)'.
FEATURES             Location/Qualifiers
     source          1. .342
                     /organism="Homo sapiens"
                     /mol_type="genomic DNA"
                     /db_xref="taxon:9606"
ORIGIN

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  Best Local Similarity  85.0%;  Pred. No. 2.8e-71;
  Matches  273;  Conservative    0;  Mismatches   48;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
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Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
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Qy        301 GGGACACGACTGGACATTCAA 321
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RESULT 15
BD097618
LOCUS       BD097618                321 bp    DNA     linear   PAT 27-AUG-2002
DEFINITION  Antibody library.
ACCESSION   BD097618
VERSION     BD097618.1  GI:22643192
KEYWORDS    WO 0162907-A/73.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 321)
  AUTHORS   Kurosawa,Y., Akahori,Y., Iba,Y., Morino,K., Shinohara,M.,
            Takahashi,M., Okuno,Y. and Shiraki,K.
  TITLE     Antibody library
  JOURNAL   Patent: WO 0162907-A 73 30-AUG-2001;
            MEDICAL & BIOLOGICAL LABORATORIES CO LTD,YOSHIKAZU KUROSAWA,
            YASUSHI AKAHORI,YOSHITAKA IBA,KAZUHIKO MORINO,MIDORI SHINOHARA,
            MOTOHIDE TAKAHASHI,YOSHINOBU OKUNO,KIMIYASU SHIRAKI
COMMENT     OS   Homo sapiens (human)
            PN   WO 0162907-A/73
            PD   30-AUG-2001
            PF   22-FEB-2001 WO 2001JP001298
            PR   22-FEB-2000 JP 00P    50543
            PI   YOSHIKAZU KUROSAWA,YASUSHI AKAHORI,YOSHITAKA IBA,KAZUHIKO  PI
             MORINO,
            PI   MIDORI SHINOHARA,MOTOHIDE TAKAHASHI,YOSHINOBU OKUNO,KIMIYASU
            PI   SHIRAKI
            PC   C12N15/09,C07K16/00//C12P21/08
            CC   Antibody library
            FH   Key             Location/Qualifiers
            FT   source          1. .321
            FT                   /organism='Homo sapiens (human)'.
FEATURES             Location/Qualifiers
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                     /mol_type="genomic DNA"
                     /db_xref="taxon:9606"
ORIGIN

  Query Match             75.6%;  Score 242.6;  DB 6;  Length 321;
  Best Local Similarity   84.7%;  Pred. No. 9.7e-71;
  Matches  272;  Conservative    0;  Mismatches   49;  Indels    0;  Gaps    0;

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Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||||| || || ||    ||||| |    | ||   ||||||||
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Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||||  | |||||   || ||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCCATCACCTTCGGCCAA 300

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Search completed: September 11, 2005, 21:22:04
Job time : 1633.32 secs
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: September 11, 2005, 17:46:48 ; Search time 337.536 Seconds
(without alignments)
5629.725 Million cell updates/sec

Title: US-09-403-107-147
Perfect score: 321
Sequence: 1 gagctccagatgacccagtc.....ggacacgactggacattcaa 321

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 4390206 seqs, 2959870667 residues

Total number of hits satisfying chosen parameters: 8780412

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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13: geneseqn2004bs:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	321	100.0	321	2	AAV68539	Aav68539 Nucleotid
2	321	100.0	1630	3	AAZ50588	Aaz50588 HD70scFv-
3	321	100.0	1630	3	AAZ50587	Aaz50587 HD70scFv-
4	260.2	81.1	321	2	AAV68536	Aav68536 Nucleotid
5	260.2	81.1	321	2	AAX77237	Aax77237 Human kap
6	249	77.6	321	11	ADO51603	Ado51603 Human TAG
7	245.8	76.6	324	10	AAL52122	Aal52122 Human ant
8	244.2	76.1	342	4	AAH47727	Aah47727 Nucleotid
9	242.6	75.6	321	4	AAH47731	Aah47731 Nucleotid
10	240.8	75.0	321	5	AAH68651	Aah68651 Human ant
11	240.8	75.0	321	9	ACD45315	Acd45315 Anti-Rh(D
12	240.2	74.8	321	12	ADO36505	Ado36505 Human ant
13	240.2	74.8	321	12	ADO36497	Ado36497 Human ant
14	238.6	74.3	327	10	ADD89879	Add89879 Human ant
15	238.6	74.3	327	13	ADS64656	Ads64656 Human 9C1
16	238.6	74.3	327	13	ADS64655	Ads64655 Human 9C1
17	237.8	74.1	321	12	ADP22249	Adp22249 Human ant
18	237.8	74.1	324	4	AAS03434	Aas03434 DNA encod
19	237.8	74.1	735	13	ADR28103	Adr28103 Polynucle
20	236.2	73.6	321	11	ADO51602	Ado51602 Human TAG
21	236.2	73.6	321	12	ADP22235	Adp22235 Human ant
22	236.2	73.6	321	12	ADP22253	Adp22253 Human ant
23	236.2	73.6	458	12	ADK52349	Adk52349 Human ant
24	235	73.2	1106	6	ABQ54241	Abq54241 Human ova
25	234.4	73.0	936	3	AAA27390	Aaa27390 Human IGF
26	233	72.6	321	12	ADP22243	Adp22243 Human ant
27	232.8	72.5	917	3	AAA27381	Aaa27381 Human IGF
28	231.4	72.1	333	5	AAH74684	Aah74684 Nucleotid
29	231.4	72.1	333	10	ABT34320	Abt34320 Hepatitis
30	231.4	72.1	720	2	AAX36070	Aax36070 DNA encod
31	231.4	72.1	720	10	ABZ76706	Abz76706 Human ser
32	231.4	72.1	720	12	ADL92368	Adl92368 Human pha
33	231.4	72.1	720	12	ADQ77166	Adq77166 HSA Heavy
34	231.4	72.1	900	5	AAH74688	Aah74688 Nucleotid
35	231.4	72.1	900	10	ABT34324	Abt34324 Hepatitis
36	229.8	71.6	321	9	AAD57375	Aad57375 Human 5F1
37	229.8	71.6	324	10	AAL52120	Aal52120 Human ant
38	229.8	71.6	396	2	AAT75423	Aat75423 Human ant
39	229.8	71.6	684	4	AAH30052	Aah30052 TRO005 ka
40	229.8	71.6	720	10	ABT34315	Abt34315 Hepatitis
41	229.8	71.6	729	3	AAA11630	Aaa11630 Human imm
42	229.8	71.6	729	6	ABL46009	Abl46009 Humanised
43	228.2	71.1	321	11	ADO51601	Ado51601 Human TAG
44	228.2	71.1	324	4	AAF29086	Aaf29086 Human HIV
45	228.2	71.1	324	4	AAF29073	Aaf29073 Human HIV

ALIGNMENTS

RESULT 1
AAV68539
ID AAV68539 standard; DNA; 321 BP.
XX
AC AAV68539;
XX
DT 16-FEB-1999 (first entry)
XX
DE Nucleotide sequence of human kappa 5.1 light chain variable region.
XX
KW Human; kappa 5.1 light chain variable region; receptor; antigen; tumour;
KW auto-immune disease; graft rejection; allergy; inflammatory disease;
KW endocrine disease; degenerative disease; ss.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT CDS 1. .321
FT /*tag= a
FT /product= "human kappa 5.1 light chain variable region"
XX
PN WO9846645-A2.
XX
PD 22-OCT-1998.
XX
PF 14-APR-1998; 98WO-EP002180.
XX
PR 14-APR-1997; 97EP-00106109.
XX
PA (KUFE/) KUFER P.
PA (RAUM/) RAUM T.
XX
PI Kufer P, Raum T;
XX
DR WPI; 1998-594564/50.
DR P-PSDB; AAW68539.
XX
PT Production of anti-human antigen receptors - by selecting a combination
PT of functionally rearranged VH and VL immunoglobulin chains expressed from
PT a recombinant vector.
XX
PS Claim 9; Fig 9; 84pp; English.
XX

```
CC   This is the nucleotide sequence of the human kappa light chain variable
CC   region, used in the method of the invention. for providing receptors that
CC   can be used for targeting antigens in humans without being immunogenic
CC   themselves. Such receptors can be used for treating diseases such as
CC   tumours or auto-immune diseases, graft rejection after transplantation,
CC   infectious diseases by targeting cellular receptors as well as allergic,
CC   inflammatory, endocrine and degenerative diseases by targeting key
CC   molecules involved in the pathological process
XX
SQ   Sequence 321 BP; 81 A; 93 C; 72 G; 75 T; 0 U; 0 Other;

  Query Match             100.0%;  Score 321;  DB 2;  Length 321;
  Best Local Similarity   100.0%;  Pred. No. 1.5e-93;
  Matches  321;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
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Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Db        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||||||||
Db        301 GGGACACGACTGGACATTCAA 321


RESULT 2
AAZ50588
ID   AAZ50588 standard; DNA; 1630 BP.
XX
AC   AAZ50588;
XX
DT   23-MAY-2000  (first entry)
XX
DE   HD70scFv-Ck-interleukin 2 encoding DNA.
XX
KW   HD70; single-chain variable fragment; scFv; 17-1A antigen; human; EpCAM;
KW   epithelial cell adhesion molecule; inflammatory cytokine; IL-2;
KW   interleukin-2; Ck-domain; kappa light chain constant domain;
KW   heterominibody; multifunctional compound; immunoglobulin; cytostatic;
KW   immunostimulatory; antileukaemia; diagnosis; prevention;
KW   antiproliferative; treatment; malignant; haematopoietic cell; lymphoma;
KW   leukaemia; solid tumour; carcinoma; melanoma; sarcoma; ds.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
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FT                   /*tag=  a
FT                   /product= "HD70scFv-Ck-IL-2 chain"
FT   misc_feature    96. .842
FT                   /*tag=  b
FT                   /label= HD70_scFv
XX
PN   WO200006605-A2.
XX
PD   10-FEB-2000.
XX
PF   28-JUL-1999;   99WO-EP005416.
XX
PR   28-JUL-1998;   98EP-00114082.
XX
PA   (MICR-) MICROMET GES BIOMEDIZINISCHE FORSCHUNG.
XX
PI   Kufer P,  Dreier T,  Baeuerle PA,  Borschert K,  Zettl F;
XX
DR   WPI; 2000-195265/17.
DR   P-PSDB; AAY44995.
XX
PT   New multifunctional compounds useful for preventing and/or treating
PT   malignant cell growth and for detection and diagnosis.
XX
PS   Claim 8; Fig 55B; 166pp; English.
XX
CC   The patent discloses heterominibodies which are multifunctional compounds
CC   produceable in a mammalian host cell as a secretable and fully functional
CC   heterodimer of two polypeptide chains, where one of the polypeptide
CC   chains comprises, a CH1-domain (constant domain of an immunoglobulin
CC   heavy chain) and the other chain comprises CL-domain (constant domain of
CC   an immunoglobulin light chain). The polypeptide chains further comprise,
CC   fused to the constant domains at least two (poly)peptides having
CC   different receptor or ligand functions, where further at least two of the
CC   different (poly)peptides lack an intrinsic affinity for one another and
CC   are linked via the constant domains. The heterominibodies have
CC   cytostatic, immunostimulatory, antileukaemia and antiproliferative
CC   activities. These compounds can be used for diagnosing, preventing and
CC   treating malignant cell growth related to malignancies of haematopoietic
CC   cells e.g. lymphomas and leukaemias, or to solid tumours e.g. carcinomas,
CC   melanomas and sarcomas. The present sequence is a DNA encoding right
CC   chain of a heterominibody comprising HD70 single-chain Fv (scFv) fragment
CC   N-terminally linked to human Ck domain (constant domain of immunoglobulin
CC   -kappa light chain) which bears at its C-terminus the human inflammatory
CC   cytokine interleukin-2 (IL-2). HD70 scFv specifically recognises the
CC   human epithelial cell adhesion molecule (EpCAM) also called 17-1A antigen
XX
SQ   Sequence 1630 BP; 445 A; 410 C; 408 G; 367 T; 0 U; 0 Other;

  Query Match             100.0%;  Score 321;  DB 3;  Length 1630;
  Best Local Similarity   100.0%;  Pred. No. 2.8e-93;
  Matches  321;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Db         96 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 155

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        156 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 215

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
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Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
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Db        336 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 395

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||||||||
Db        396 GGGACACGACTGGACATTCAA 416


RESULT 3
AAZ50587
ID   AAZ50587 standard; DNA; 1630 BP.
XX
AC   AAZ50587;
XX
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DT   23-MAY-2000  (first entry)
XX
DE   HD70scFv-CH1-GM-CSF chain encoding DNA.
XX
KW   HD70; single-chain variable fragment; scFv; 17-1A antigen; human; EpCAM;
KW   epithelial cell adhesion molecule; inflammatory cytokine; GM-CSF;
KW   granulocyte/macrophage colony stimulating factor; heterominibody;
KW   CH1-domain; multifunctional compound; heavy chain constant domain;
KW   immunoglobulin; cytostatic; immunostimulatory; antileukaemia; diagnosis;
KW   antiproliferative; prevention; treatment; malignant; haematopoietic cell;
KW   lymphoma; leukaemia; solid tumour; carcinoma; melanoma; sarcoma; ds.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
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FT   misc_feature    96. .842
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PN   WO200006605-A2.
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PD   10-FEB-2000.
XX
PF   28-JUL-1999;   99WO-EP005416.
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PR   28-JUL-1998;   98EP-00114082.
XX
PA   (MICR-) MICROMET GES BIOMEDIZINISCHE FORSCHUNG.
XX
PI   Kufer P,  Dreier T,  Baeuerle PA,  Borschert K,  Zettl F;
XX
DR   WPI; 2000-195265/17.
DR   P-PSDB; AAY44994.
XX
PT   New multifunctional compounds useful for preventing and/or treating
PT   malignant cell growth and for detection and diagnosis.
XX
PS   Claim 8; Fig 55A; 166pp; English.
XX
CC   The patent discloses heterominibodies which are multifunctional compounds
CC   produceable in a mammalian host cell as a secretable and fully functional
CC   heterodimer of two polypeptide chains, where one of the polypeptide
CC   chains comprises, a CH1-domain (constant domain of an immunoglobulin
CC   heavy chain) and the other chain comprises CL-domain (constant domain of
CC   an immunoglobulin light chain). The polypeptide chains further comprise,
CC   fused to the constant domains at least two (poly)peptides having
CC   different receptor or ligand functions, where further at least two of the
CC   different (poly)peptides lack an intrinsic affinity for one another and
CC   are linked via the constant domains. The heterominibodies have
CC   cytostatic, immunostimulatory, antileukaemia and antiproliferative
CC   activities. These compounds can be used for diagnosing, preventing and
CC   treating malignant cell growth related to malignancies of haematopoietic
CC   cells e.g. lymphomas and leukaemias, or to solid tumours e.g. carcinomas,
CC   melanomas and sarcomas. The present sequence is a DNA encoding left chain
CC   of a heterominibody comprising HD70 single-chain Fv (scFv) fragment N-
CC   terminally linked to human CH1 domain which bears at its C-terminus the
CC   human inflammatory cytokine granulocyte/macrophage colony stimulating
CC   factor (GM-CSF), plus a hexahistidine sequence for ease of purification.
CC   HD70 scFv specifically recognises the human epithelial cell adhesion
CC   molecule (EpCAM) also called 17-1A antigen
XX
SQ   Sequence 1630 BP; 376 A; 484 C; 437 G; 333 T; 0 U; 0 Other;

  Query Match             100.0%;  Score 321;  DB 3;  Length 1630;
  Best Local Similarity   100.0%;  Pred. No. 2.8e-93;
  Matches  321;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         96 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 155

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
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Db        336 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 395

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||||||||
Db        396 GGGACACGACTGGACATTCAA 416


RESULT 4
AAV68536
ID   AAV68536 standard; DNA; 321 BP.
XX
AC   AAV68536;
XX
DT   16-FEB-1999  (first entry)
XX
DE   Nucleotide sequence of human Kappa 8 light chain variable region.
XX
KW   Human; kappa 8 light chain variable region; receptor; antigen; tumour;
KW   auto-immune disease; graft rejection; allergy; inflammatory disease;
KW   endocrine disease; degenerative disease; ss.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   CDS             1. .321
FT                   /*tag=  a
FT                   /product= "human kappa 8 light chain variable region"
XX
PN   WO9846645-A2.
XX
PD   22-OCT-1998.
XX
PF   14-APR-1998;   98WO-EP002180.
XX
PR   14-APR-1997;   97EP-00106109.
XX
PA   (KUFE/) KUFER P.
PA   (RAUM/) RAUM T.
XX
PI   Kufer P,  Raum T;
XX
DR   WPI; 1998-594564/50.
DR   P-PSDB; AAW80814.
XX
PT   Production of anti-human antigen receptors - by selecting a combination
PT   of functionally rearranged VH and VL immunoglobulin chains expressed from
PT   a recombinant vector.
XX
PS   Claim 9; Fig 6; 84pp; English.
XX
CC   This is the nucleotide sequence of the human kappa 8 light chain variable
CC   region, used in the method of the invention. for providing receptors that
CC   can be used for targeting antigens in humans without being immunogenic
CC   themselves. Such receptors can be used for treating diseases such as
CC   tumours or auto-immune diseases, graft rejection after transplantation,
CC   infectious diseases by targeting cellular receptors as well as allergic,
CC   inflammatory, endocrine and degenerative diseases by targeting key
CC   molecules involved in the pathological process
```

```
XX
SQ   Sequence 321 BP; 81 A; 91 C; 75 G; 74 T; 0 U; 0 Other;

  Query Match             81.1%;  Score 260.2;  DB 2;  Length 321;
  Best Local Similarity   88.2%;  Pred. No. 7.5e-74;
  Matches  283;  Conservative    0;  Mismatches   38;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              |||||||||||||||||||||||||||||||||||| ||||| ||||||||||||||||
Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCTTCTGTGGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||| ||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGTCGGACAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              |||||||| ||||| |  ||||||||| ||| ||||||||||||||||| || || |||
Db        181 CGATTCAGTGGCAGCGGGTCTGGGACAGATTTCACTCTCACCATCAGCAGTCTACAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||| ||| |||||||  ||||| |||  | ||    | |||  ||| || |||||
Db        241 GAAGATTCTGCAACTTACTACTGTCAGCAGAGTTACGACATCCCGTACACTTTTGGCCAG 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||    ||||| ||  ||
Db        301 GGGACCAAGCTGGAGATCAAA 321


RESULT 5
AAX77237
ID   AAX77237 standard; DNA; 321 BP.
XX
AC   AAX77237;
XX
DT   04-AUG-1999  (first entry)
XX
DE   Human kappa 8 light chain variable region encoding DNA.
XX
KW   Binding site domain; BSD; epitope; fusion protein; therapeutic; cancer;
KW   autoimmune disease; scFv-antibody; single-chain Fv; ss.
XX
OS   Homo sapiens.
XX
PN   WO9925818-A1.
XX
PD   27-MAY-1999.
XX
PF   16-NOV-1998;   98WO-EP007313.
XX
PR   17-NOV-1997;   97EP-00120096.
XX
PA   (KUFE/) KUFER P.
XX
PI   Kufer P,  Raum T,  Borschert K,  Zettl F,  Lutterbuese R;
XX
DR   WPI; 1999-338004/28.
DR   P-PSDB; AAY17955.
XX
PT   Phage display system for identification of binding site domains retaining
PT   capacity to bind an epitope.
XX
PS   Disclosure; Fig 3.2; 152pp; English.
XX
CC   The invention relates to a method of identifying binding site domains
CC   (BSD) that retain the capacity of binding to a predetermined epitope when
CC   positioned C-terminal of at least one further domain in a recombinant bi-
CC   or multivalent polypeptide. The method comprises (a) testing a panel of
CC   BSD displayed on the surface of a biological display system as part of a
CC   fusion protein for binding to a predetermined epitope, where the fusion
CC   protein comprises an additional domain positioned N-terminal of the BSD
CC   and an amino acid sequence that mediates anchoring of the fusion protein
CC   to the surface of the display system; and (b) identifying a BSD that
CC   binds to the predetermined epitope. The method is useful to identify bi-
CC   or multivalent polypeptides that comprise antibody binding sites capable
CC   of efficiently binding to the corresponding antigen. The polypeptides or
CC   antibodies identified by the method are useful therapeutically and
CC   diagnostically, for e.g. cancer and autoimmune diseases. ScFv-antibody
CC   fragments that bind independently of their position within bifunctional
CC   single-chain fusion proteins can be isolated from combinatorial antibody
CC   libraries using the new in vitro method
XX
SQ   Sequence 321 BP; 81 A; 91 C; 75 G; 74 T; 0 U; 0 Other;

  Query Match             81.1%;  Score 260.2;  DB 2;  Length 321;
  Best Local Similarity   88.2%;  Pred. No. 7.5e-74;
  Matches  283;  Conservative    0;  Mismatches   38;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              |||||||||||||||||||||||||||||||||||| ||||| ||||||||||||||||
Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCTTCTGTGGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||| ||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGTCGGACAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              |||||||| ||||| |  ||||||||| ||| ||||||||||||||||| || || |||
Db        181 CGATTCAGTGGCAGCGGGTCTGGGACAGATTTCACTCTCACCATCAGCAGTCTACAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||| ||| ||||||||  ||||| |||  | ||    | |||  ||| || |||||
Db        241 GAAGATTCTGCAACTTACTACTGTCAGCAGAGTTACGACATCCCGTACACTTTTGGCCAG 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||    ||||| ||  ||
Db        301 GGGACCAAGCTGGAGATCAAA 321


RESULT 6
ADO51603
ID   ADO51603 standard; DNA; 321 BP.
XX
AC   ADO51603;
XX
DT   15-JUL-2004  (first entry)
XX
DE   Human TAG-72 antibody-related KCS18 gene SeqID13.
XX
KW   semi-human monoclonal antibody; tumour-associated glycoprotein antigen;
KW   TAG-72; human light chain; cancer; AKA; HzK; VK1;
KW   human immunoglobulin kappa light chain germline; gene; ds; human.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   CDS             1. .321
FT                   /*tag=  a
FT                   /product= "Human TAG-72 antibody-related KCS18 protein"
FT                   /partial
FT                   /note= "No start or stop codon"
XX
PN   KR2003013633-A.
XX
PD   15-FEB-2003.
XX
PF   08-AUG-2001; 2001KR-00047737.
XX
PR   08-AUG-2001; 2001KR-00047737.
```

```
XX
PA   (KORE-) KOREA RES INST BIOSCIENCE & BIOTECHNOLOG.
XX
PI   Hong HJ,  Kim SJ;
XX
DR   WPI; 2003-500992/47.
DR   P-PSDB; ADO51598.
XX
PT   Semi-human monoclonal antibody binding to tumor-associated glycoprotein
PT   antigen tag-72 and human light chain comprising the same, useful for
PT   diagnosing and treating cancer.
XX
PS   Claim 13; SEQ ID NO 18; 1pp; Korean.
XX
CC   This invention relates to a novel semi-human monoclonal antibody binding
CC   to tumour-associated glycoprotein antigen TAG-72 and a human light chain
CC   comprising the same. The semi-human monoclonal antibody has a human light
CC   chain of which amino acid sequence is completely derived from the human,
CC   therefore it is useful for the diagnosis and treatment of cancer. The
CC   semi-human monoclonal antibody specifically binds to TAG-72 antigen of
CC   which the light chain of human antibody AKA/HzK is substituted with the
CC   light chain of a human derived antibody. The light chain of the human
CC   derived antibody is preferably composed of amino acid sequence derived
CC   from the VK1 family of the human immunoglobulin kappa light chain
CC   germline. The present sequence is that of a gene which encodes a human
CC   KCS protein which is related to the novel semi-human TAG-72 antibody of
CC   the invention.
XX
SQ   Sequence 321 BP; 87 A; 86 C; 73 G; 75 T; 0 U; 0 Other;

  Query Match             77.6%;  Score 249;  DB 11;  Length 321;
  Best Local Similarity   86.0%;  Pred. No. 3.2e-70;
  Matches  276;  Conservative    0;  Mismatches   45;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |||||
Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAATCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||||||||||||| ||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGGCAAGTCAGGGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||||| || || ||   |||||||     |||   ||||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATAAGGCATCTAGTTTAGAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||||||||||| |||||||||| | | ||||||||||||||||||||||||||||||
Db        181 AGGTTCAGCGGCAGTGGATCTGGGACAGAATTCACTCTCACCATCAGCAGCCTGCAGCCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              || ||||||| |||||| |  |||||||| || ||| |   || ||||||||||||||||
Db        241 GATGATTTTGGTACTTATTACTGTCAACAGACTTACATTGCCCCCATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 7
AAL52122
ID   AAL52122 standard; DNA; 324 BP.
XX
AC   AAL52122;
XX
DT   29-MAY-2003  (first entry)
XX
DE   Human anti-blood coagulation factor VIII antibody-related gene #4.
XX
KW   Human; gene; ds; anti-blood coagulation factor VIII antibody;
KW   FVIII antibody; anti-thrombotic; thrombosis; activated FVIII.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   CDS             1. .324
FT                   /*tag= a
FT                   /partial
FT                   /product= "Human VIII antibody-related protein #4"
FT                   /note= "No start or stop codon is given"
XX
PN   WO2002101040-A1.
XX
PD   19-DEC-2002.
XX
PF   11-JUN-2002; 2002WO-JP005783.
XX
PR   12-JUN-2001; 2001JP-00177640.
XX
PA   (KAGA ) CHEMO-SERO-THERAPEUTIC RES INST.
XX
PI   Nakashima T,  Yuguchi M;
XX
DR   WPI; 2003-148804/14.
DR   P-PSDB; AAO16706.
XX
PT   Human-type anti-blood coagulation factor VIII antibody, applicable in
PT   antithrombotics for prevention or treatment of thrombosis, in diagnosis
PT   of activated FVIII, and for maintaining low coagulation state.
XX
PS   Disclosure; Page 28; 39pp; Japanese.
XX
CC   The invention comprises the amino acid and coding sequence of a human
CC   anti-blood coagulation factor VIII (FVIII) antibody, which has an
CC   inhibitory activity on the coagulation activity of human FVIII. The
CC   antibody is applicable in anti-thrombotics for prevention or treatment of
CC   thrombosis, and in the diagnosis of activated FVIII. The present DNA
CC   sequence encodes a human FVIII antibody-related protein
XX
SQ   Sequence 324 BP; 87 A; 88 C; 73 G; 76 T; 0 U; 0 Other;

  Query Match             76.6%;  Score 245.8;  DB 10;  Length 324;
  Best Local Similarity   85.4%;  Pred. No. 3.5e-69;
  Matches  274;  Conservative    0;  Mismatches   47;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||||| || || ||    |||||  |    | ||   ||||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||| | |||||   ||||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 8
AAH47727
ID   AAH47727 standard; DNA; 342 BP.
XX
AC   AAH47727;
XX
```

DT 30-NOV-2001 (first entry)
XX
DE Nucleotide sequence of seq Id No. 70.
XX
KW Gene library; immunoglobulin; antibody library; human; ds.
XX
OS Homo sapiens.
XX
PN WO200162907-A1.
XX
PD 30-AUG-2001.
XX
PF 22-FEB-2001; 2001WO-JP001298.
XX
PR 22-FEB-2000; 2000JP-00050543.
XX
PA (MEDI-) MEDICAL & BIOLOGICAL LAB CO LTD.
XX
PI Kurosawa Y, Akahori Y, Iba Y, Morino K, Shinohara M, Takahashi M;
PI Okuno Y, Shiraki K;
XX
DR WPI; 2001-565420/63.
DR P-PSDB; AAG65563.
XX
PT Producing gene libraries and antibody libraries, involves selecting a
PT light chain that binds to a heavy chain product to produce a functional
PT formation, and producing a gene library of the light chain variable
PT regions.
XX
PS Examples; p 146-147; 181pp; Japanese.
XX
CC The invention relates to producing gene libraries, comprising
CC immunoglobulin light and heavy variable region. The method involves
CC selecting light chain that binds with the heavy chain product to produce
CC a functional conformation, producing a gene library comprising a
CC collection of these light chain variable genes, and combining with gene
CC library of heavy chain variable genes. The method is used for production
CC of gene and antibody libraries
XX
SQ Sequence 342 BP; 91 A; 93 C; 78 G; 80 T; 0 U; 0 Other;

Query Match 76.1%; Score 244.2; DB 4; Length 342;
Best Local Similarity 85.0%; Pred. No. 1.2e-68;
Matches 273; Conservative 0; Mismatches 48; Indels 0; Gaps 0;

```
Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCGTGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||||| || || ||    ||||| |    | ||   |||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| ||||||| |||||||||| ||| ||||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||  | |||||  |||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321
```

RESULT 9
AAH47731
ID AAH47731 standard; DNA; 321 BP.
XX
AC AAH47731;
XX
DT 30-NOV-2001 (first entry)
XX
DE Nucleotide sequence of seq Id No. 74.
XX
KW Gene library; immunoglobulin; antibody library; human; ds.
XX
OS Homo sapiens.
XX
PN WO200162907-A1.
XX
PD 30-AUG-2001.
XX
PF 22-FEB-2001; 2001WO-JP001298.
XX
PR 22-FEB-2000; 2000JP-00050543.
XX
PA (MEDI-) MEDICAL & BIOLOGICAL LAB CO LTD.
XX
PI Kurosawa Y, Akahori Y, Iba Y, Morino K, Shinohara M, Takahashi M;
PI Okuno Y, Shiraki K;
XX
DR WPI; 2001-565420/63.
DR P-PSDB; AAG65567.
XX
PT Producing gene libraries and antibody libraries, involves selecting a
PT light chain that binds to a heavy chain product to produce a functional
PT formation, and producing a gene library of the light chain variable
PT regions.
XX
PS Examples; p 149; 181pp; Japanese.
XX
CC The invention relates to producing gene libraries, comprising
CC immunoglobulin light and heavy variable region. The method involves
CC selecting light chain that binds with the heavy chain product to produce
CC a functional conformation, producing a gene library comprising a
CC collection of these light chain variable genes, and combining with gene
CC library of heavy chain variable genes. The method is used for production
CC of gene and antibody libraries
XX
SQ Sequence 321 BP; 87 A; 87 C; 72 G; 75 T; 0 U; 0 Other;

Query Match 75.6%; Score 242.6; DB 4; Length 321;
Best Local Similarity 84.7%; Pred. No. 3.8e-68;
Matches 272; Conservative 0; Mismatches 49; Indels 0; Gaps 0;

```
Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCGTGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||||| || || ||    ||||| |    | ||   |||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| ||||||| |||||||||| ||| ||||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||  | |||||  || ||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCCATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321
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RESULT 10
AAH68651
ID   AAH68651 standard; DNA; 321 BP.
XX
AC   AAH68651;
XX
DT   14-SEP-2001  (first entry)
XX
DE   Human anti-Rh(D) chain I06 nucleotide sequence.
XX
KW   Human; anti-Rh(D) antibody; Rh(D) binding protein; immunostimulant;
KW   red blood cell; Rh phenotype; diagnosis; therapeutic; ds.
XX
OS   Homo sapiens.
XX
PN   US6255455-B1.
XX
PD   03-JUL-2001.
XX
PF   29-JAN-1999;   99US-00240274.
XX
PR   11-OCT-1996;   96US-0028550P.
PR   27-JUN-1997;   97US-00884045.
PR   10-APR-1998;   98US-0081380P.
XX
PA   (UYPE-) UNIV PENNSYLVANIA.
XX
PI   Siegel DL;
XX
DR   WPI; 2001-388931/41.
DR   P-PSDB; AAG93594.
XX
PT   New isolated protein, preferably a human anti-Rh(D) antibody for use in
PT   diagnostics requiring a human instead of an animal antibody and in
PT   therapeutic medicine.
XX
PS   Example 2; Col 55; 162pp; English.
XX
CC   The present invention describes an isolated Rh(D) binding protein,
CC   preferably a human antibody, (I) having an amino acid sequence comprising
CC   one of the sequences (S) given in AAG93558 to AAG93669. (I) has
CC   immunostimulant activity, and can be used as an immune system stimulant.
CC   (I) can be used in diagnostic and therapeutic medicine. The antibodies
CC   are used in diagnostics that require human antibodies instead of animal
CC   antibodies, such as determine the Rh phenotype of human red blood cells.
CC   AAH68615 to AAH68726 represent the nucleotide sequence which encode
CC   AAG93558 to AAG93669. AAG93670 to AAG93697 represent anti-Rh(D) heavy
CC   chain CDR3 amino acid sequences which are given in the exemplification of
CC   the present invention
XX
SQ   Sequence 321 BP; 86 A; 90 C; 73 G; 72 T; 0 U; 0 Other;

  Query Match             75.0%;  Score 240.8;  DB 5;  Length 321;
  Best Local Similarity   85.1%;  Pred. No. 1.4e-67;
  Matches  269;  Conservative    0;  Mismatches   47;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  |
Db         63 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
                | |||||||| || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAAGCTCCTGATCTATGCCGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| ||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| ||||||| ||||||||| | |||||  ||||||||||||||||||||||||
Db        243 TTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAAGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
              |||||||||  ||| ||
Db        303 ACGACTGGAGATTAAA 318


RESULT 11
ACD45315
ID   ACD45315 standard; DNA; 321 BP.
XX
AC   ACD45315;
XX
DT   12-SEP-2003  (first entry)
XX
DE   Anti-Rh(D) chain I06 DNA.
XX
KW   Human; ds; gene; RH(D) binding protein; blood typing; blood product;
KW   magnetically activated cell sorting.
XX
OS   Homo sapiens.
XX
PN   US2003040605-A1.
XX
PD   27-FEB-2003.
XX
PF   04-MAY-2001; 2001US-00848798.
XX
PR   11-OCT-1996;   96US-0028550P.
PR   27-JUN-1997;   97US-00884045.
PR   10-APR-1998;   98US-0081380P.
PR   29-JAN-1999;   99US-00240274.
XX
PA   (UYPE-) UNIV PENNSYLVANIA.
XX
PI   Siegel DL;
XX
DR   WPI; 2003-512273/48.
DR   P-PSDB; ABO27401.
XX
PT   New human Rh(D)-binding protein useful for various diagnostic and
PT   therapeutic applications, including typing of blood or blood products.
XX
PS   Claim 12; Page 40; 187pp; English.
XX
CC   The invention relates to an isolated Rh(D) binding protein. The protein
CC   can be used for magnetically activated cell sorting. The protein is
CC   useful in various diagnostic and therapeutic applications in humans,
CC   including typing of blood or blood products. The present sequence
CC   represents DNA encoding a human anti-Rh(D) chain
XX
SQ   Sequence 321 BP; 86 A; 90 C; 73 G; 72 T; 0 U; 0 Other;

  Query Match             75.0%;  Score 240.8;  DB 9;  Length 321;
  Best Local Similarity   85.1%;  Pred. No. 1.4e-67;
  Matches  269;  Conservative    0;  Mismatches   47;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  |
Db         63 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
                | |||||||| || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAAGCTCCTGATCTATGCCGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| ||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242
```

```
Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| |||||||  |||||||||  | |||||   ||||||||||||||||||||||||
Db        243 TTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAAGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
              ||||||||| ||| ||
Db        303 ACGACTGGAGATTAAA 318
```

RESULT 12
ADO36505
ID ADO36505 standard; cDNA; 321 BP.
XX
AC ADO36505;
XX
DT 12-AUG-2004 (first entry)
XX
DE Human anti-heparanase IE7 Vk encoding cDNA SEQ ID NO:19.
XX
KW human; monoclonal antibody; heparanase; heparanase inhibitor;
KW anti-heparanase antibody; cytostatic; immunosuppressive; antiarthritic;
KW antiasthmatic; antiinflammatory; dermatological; antiarteriosclerotic;
KW neuroprotective; nootropic; heparanase antagonist; cancer; tumour;
KW melanoma; lymphoma; prostate carcinoma; pancreatic carcinoma;
KW bladder carcinoma; fibrosarcoma; rhabdomyosarcoma; mastocytoma;
KW mammary adenocarcinoma; leukaemia; rheumatoid fibroblast;
KW autoimmune disease; arthritis; asthma; lupus erythematosus;
KW allograft rejection; vascular restenosis; atherosclerosis;
KW Alzheimer's disease; gene; ss.
XX
OS Homo sapiens.
XX
FH Key   Location/Qualifiers
FT CDS   1. .321
FT   /*tag= a
FT   /product= "anti-heparanase IE7 Vk"
XX
PN WO2004043989-A2.
XX
PD 27-MAY-2004.
XX
PF 05-NOV-2003; 2003WO-US035464.
XX
PR 07-NOV-2002; 2002US-0424803P.
XX
PA (MEDA-) MEDAREX INC.
PA (CELL-) CELLTECH R & D.
XX
PI Huang H, Holmes S, Mason S;
XX
DR WPI; 2004-411694/38.
DR P-PSDB; ADO36506.
XX
PT New human monoclonal antibody to heparanase, for use in treating or
PT preventing cancer, autoimmune disease, arthritis, asthma, lupus
PT erythematosus, allograft rejection, atherosclerosis, and Alzheimer's
PT disease.
XX
PS Claim 8; SEQ ID NO 19; 108pp; English.
XX
CC The present invention describes an isolated human monoclonal antibody
CC which binds to and inhibits activity of human heparanase. Human anti-
CC heparanase antibodies of the present invention have cytostatic,
CC immunosuppressive, antiarthritic, antiasthmatic, antiinflammatory,
CC dermatological, antiarteriosclerotic, neuroprotective and nootropic
CC activities, and can be used as heparanase antagonists. The antibody,
CC methods and compositions of the present invention are useful in treating
CC or preventing cancer or tumours, e.g. melanoma, lymphoma, prostate
CC carcinoma, pancreatic carcinoma, bladder carcinoma, fibrosarcoma,
CC rhabdomyosarcoma, mastocytoma, mammary adenocarcinoma, leukaemia or a
CC rheumatoid fibroblast, autoimmune disease, arthritis, asthma, lupus
CC erythematosus, allograft rejection, vascular restenosis, atherosclerosis,
CC and Alzheimer's disease. The present sequence encodes a human anti-
CC heparanase IE7 Vk amino acid sequence, which is used in the
CC exemplification of the present invention.
XX
SQ Sequence 321 BP; 81 A; 87 C; 76 G; 77 T; 0 U; 0 Other;

```
  Query Match             74.8%;  Score 240.2;  DB 12;  Length 321;
  Best Local Similarity   84.9%;  Pred. No. 2.3e-67;
  Matches  269;  Conservative    0;  Mismatches   48;  Indels    0;  Gaps    0;

Qy          5 TCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64
              ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          5 TCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64

Qy         65 CTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGAC 124
              |||||||||||||||| ||||||||||    ||||   ||||||||||||||||||||
Db         65 CTTGCCGGGCAAGTCAGGGCATTAGCAGTGCTTTAGCCTGGTATCAGCAGAAACCAGGGA 124

Qy        125 AGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGAT 184
              |  ||||||||||  || || ||     || || |    ||||   ||||||||     | |
Db        125 AAGCTCCTAAGCTCCTGATCTATGATGCCTCCAGTTTGGAAAGTGGGGTCCCATCAAGGT 184

Qy        185 TCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244
              |||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||||
Db        185 TCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244

Qy        245 ATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGA 304
              ||||||| ||||| |  |||||||||| | | ||||  ||||||||||||||||||||
Db        245 ATTTTGCAACTTATTACTGTCAACAGTTTAATAGTTACCCGATCACCTTCGGCCAAGGGA 304

Qy        305 CACGACTGGACATTCAA 321
              |||||||||| ||| ||
Db        305 CACGACTGGAGATTAAA 321
```

RESULT 13
ADO36497
ID ADO36497 standard; cDNA; 321 BP.
XX
AC ADO36497;
XX
DT 12-AUG-2004 (first entry)
XX
DE Human anti-heparanase 13B2 VL encoding cDNA SEQ ID NO:11.
XX
KW human; monoclonal antibody; heparanase; heparanase inhibitor;
KW anti-heparanase antibody; cytostatic; immunosuppressive; antiarthritic;
KW antiasthmatic; antiinflammatory; dermatological; antiarteriosclerotic;
KW neuroprotective; nootropic; heparanase antagonist; cancer; tumour;
KW melanoma; lymphoma; prostate carcinoma; pancreatic carcinoma;
KW bladder carcinoma; fibrosarcoma; rhabdomyosarcoma; mastocytoma;
KW mammary adenocarcinoma; leukaemia; rheumatoid fibroblast;
KW autoimmune disease; arthritis; asthma; lupus erythematosus;
KW allograft rejection; vascular restenosis; atherosclerosis;
KW Alzheimer's disease; gene; ss.
XX
OS Homo sapiens.
XX
FH Key   Location/Qualifiers
FT CDS   1. .321
FT   /*tag= a
FT   /product= "anti-heparanase 13B2 VL"
XX
PN WO2004043989-A2.
XX
PD 27-MAY-2004.
XX
PF 05-NOV-2003; 2003WO-US035464.
XX
PR 07-NOV-2002; 2002US-0424803P.
XX
PA (MEDA-) MEDAREX INC.
PA (CELL-) CELLTECH R & D.

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XX
PI   Huang H,  Holmes S,  Mason S;
XX
DR   WPI; 2004-411694/38.
DR   P-PSDB; ADO36498.
XX
PT   New human monoclonal antibody to heparanase, for use in treating or
PT   preventing cancer, autoimmune disease, arthritis, asthma, lupus
PT   erythematosus, allograft rejection, atherosclerosis, and Alzheimer's
PT   disease.
XX
PS   Claim 8; SEQ ID NO 11; 108pp; English.
XX
CC   The present invention describes an isolated human monoclonal antibody
CC   which binds to and inhibits activity of human heparanase. Human anti-
CC   heparanase antibodies of the present invention have cytostatic,
CC   immunosuppressive, antiarthritic, antiasthmatic, antiinflammatory,
CC   dermatological, antiarteriosclerotic, neuroprotective and nootropic
CC   activities, and can be used as heparanase antagonists. The antibody,
CC   methods and compositions of the present invention are useful in treating
CC   or preventing cancer or tumours, e.g. melanoma, lymphoma, prostate
CC   carcinoma, pancreatic carcinoma, bladder carcinoma, fibrosarcoma,
CC   rhabdomyosarcoma, mastocytoma, mammary adenocarcinoma, leukaemia or a
CC   rheumatoid fibroblast, autoimmune disease, arthritis, asthma, lupus
CC   erythematosus, allograft rejection, vascular restenosis, atherosclerosis,
CC   and Alzheimer's disease. The present sequence encodes a human anti-
CC   heparanase 13B2 VL amino acid sequence, which is used in the
CC   exemplification of the present invention.
XX
SQ   Sequence 321 BP; 81 A; 87 C; 76 G; 77 T; 0 U; 0 Other;

  Query Match             74.8%;  Score 240.2;  DB 12;  Length 321;
  Best Local Similarity   84.9%;  Pred. No. 2.3e-67;
  Matches  269;  Conservative    0;  Mismatches   48;  Indels    0;  Gaps    0;

Qy          5 TCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64
              ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          5 TCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64

Qy         65 CTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGAC 124
              |||||||||||||||| ||||||||||   ||||   ||||||||||||||||||||
Db         65 CTTGCCGGGCAAGTCAGGGCATTAGCAGTGCTTTAGCCTGGTATCAGCAGAAACCAGGGA 124

Qy        125 AGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGAT 184
              |  |||||||||| || || ||    || || |    ||||   ||||||||     | |
Db        125 AAGCTCCTAAGCTCCTGATCTATGATGCCTCCAGTTTGGAAAGTGGGGTCCCATCAAGGT 184

Qy        185 TCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244
              |||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||||
Db        185 TCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244

Qy        245 ATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGA 304
              ||||||| ||||| |  |||||||||| | | ||||  ||||||||||||||||||||
Db        245 ATTTTGCAACTTATTACTGTCAACAGTTTAATAGTTACCCGATCACCTTCGGCCAAGGGA 304

Qy        305 CACGACTGGACATTCAA 321
              |||||||||| ||| ||
Db        305 CACGACTGGAGATTAAA 321


RESULT 14
ADD89879
ID   ADD89879 standard; cDNA; 327 BP.
XX
AC   ADD89879;
XX
DT   29-JAN-2004  (first entry)
XX
DE   Human anti-TNF antibody 9C1A light chain variable region coding sequence.
XX
KW   Human; Tumour necrosis factor; TNF; antibody; cytostatic; anabolic;
KW   eating-disorders-gen; immunomodulator; antimicrobial; cardiovascular-gen;
KW   neuroprotective; gene; ss.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   CDS             1. .327
FT                   /*tag=  a
FT                   /partial
FT                   /product= "9C1A light chain variable region"
FT                   /note= "No start or stop codon"
XX
PN   WO2003083061-A2.
XX
PD   09-OCT-2003.
XX
PF   24-MAR-2003; 2003WO-US009072.
XX
PR   26-MAR-2002; 2002US-0367903P.
XX
PA   (CENZ ) CENTOCOR INC.
XX
PI   Giles-Komar J,  Scallon BJ,  Carton JM;
XX
DR   WPI; 2003-804040/75.
DR   P-PSDB; ADD89870.
XX
PT   New isolated mammalian anti-tumor necrosis factor (TNF) antibody, useful
PT   for diagnosing or treating an anti-TNF related condition, e.g. cancer,
PT   anorexia, cachexia, or bacterial infection.
XX
PS   Example 4; Fig 3A; 87pp; English.
XX
CC   The present sequence is the coding sequence of the light chain variable
CC   region of human anti-tumour necrosis factor (TNF) monoclonal antibody
CC   9C1A. This human TNF reactive IgG monoclonal antibody was generated by
CC   cloning variable and constant region DNA in vector pC4 and expression in
CC   CHO cells. The invention provides isolated human, primate, rodent,
CC   mammalian, chimeric, humanized and/or CDR-grafted anti-TNF antibodies,
CC   immunoglobulins, their cleavage products, other specified portions and
CC   variants, as well as anti-TNF antibody compositions, nucleic acids
CC   encoding these, vectors, host cells, methods for producing the antibodies
CC   using a host cell, transgenic animal or transgenic plant or plant cell,
CC   and therapeutic compositions, methods and devices. The antibody, nucleic
CC   acid, protein, composition and methods are useful for diagnosing or
CC   treating an anti-TNF related condition, e.g. cancer, anorexia, cachexia,
CC   or an immune, cardiovascular, infectious, and/or neurological disease.
XX
SQ   Sequence 327 BP; 84 A; 89 C; 76 G; 78 T; 0 U; 0 Other;

  Query Match             74.3%;  Score 238.6;  DB 10;  Length 327;
  Best Local Similarity   84.5%;  Pred. No. 7.5e-67;
  Matches  268;  Conservative    0;  Mismatches   49;  Indels    0;  Gaps    0;

Qy          5 TCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64
              ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          5 TCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64

Qy         65 CTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGAC 124
              |||||||||||||||| ||||||||||   ||||   ||||||||||||||||||||
Db         65 CTTGCCGGGCAAGTCAGGGCATTAGCAGTGCTTTAGCCTGGTATCAGCAGAAACCAGGGA 124

Qy        125 AGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGAT 184
              |  |||||||||| || || ||    || ||       ||||   ||||||||     | |
Db        125 AAGCTCCTAAGCTCCTGATCTATGATGCCTCCTATTTGGAAAGTGGGGTCCCATCAAGGT 184

Qy        185 TCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244
              |||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||||
Db        185 TCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244

Qy        245 ATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGA 304
              ||||||| ||||| |  |||||||||| | | ||||  ||||||||||||||||||||
Db        245 ATTTTGCAACTTATTACTGTCAACAGTTTAAAAGTTACCCGATCACCTTCGGCCAAGGGA 304

Qy        305 CACGACTGGACATTCAA 321
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                ||||||||||| ||| ||
Db        305 CACGACTGGAGATTAAA 321


RESULT 15
ADS64656
ID   ADS64656 standard; DNA; 327 BP.
XX
AC   ADS64656;
XX
DT   16-DEC-2004  (first entry)
XX
DE   Human 9C1LC light chain protein encoding DNA.
XX
KW   Tumour necrosis factor; TNF; immunotherapy; TNF related diseases;
KW   obesity; immune related disease; rheumatoid arthritis;
KW   cardiovascular disease; stroke; malignant disease; leukaemia;
KW   neurological disease; multiple sclerosis; infection; hepatitis;
KW   anorectic; antiarthritic; cerebroprotective; vasotropic; cytostatic;
KW   neuroprotective; antibacterial; antiinflammatory; hepatotropic; virucide;
KW   human; 9C1LC; light chain; gene; ds.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   CDS             10. .327
FT                   /*tag=  a
FT                   /partial
FT                   /note= "No start and stop codon"
FT                   /product= "9C1LC light chain protein"
XX
PN   US2004185047-A1.
XX
PD   23-SEP-2004.
XX
PF   21-MAR-2003; 2003US-00394471.
XX
PR   21-MAR-2003; 2003US-00394471.
XX
PA   (GILE/) GILES-KOMAR J.
PA   (SCAL/) SCALLON B J.
PA   (CART/) CARTON J M.
XX
PI   Giles-Komar J,  Scallon BJ,  Carton JM;
XX
DR   WPI; 2004-676151/66.
DR   P-PSDB; ADS64657.
XX
PT   Novel isolated mammalian anti-tumor necrosis factor (TNF) alpha antibody
PT   capable of inhibiting binding of TNF alpha to TNF receptor, useful for
PT   treating TNF-related diseases such as obesity or rheumatoid arthritis.
XX
PS   Example 4; Fig 3; 45pp; English.
XX
CC   The present invention relates to a mammalian anti-tumour necrosis factor
CC   (TNF) antibody capable of inhibiting binding of TNF alpha to TNF
CC   receptor. The invention is useful for diagnosing or treating an anti-TNF
CC   related condition in a cell, tissue, organ or animal and in
CC   immunotherapy. The invention is also useful for treating TNF related
CC   diseases chosen from obesity, immune related disease such as rheumatoid
CC   arthritis, cardiovascular disease such as stroke, malignant disease such
CC   as leukaemia, neurological disease such as multiple sclerosis and
CC   bacterial or viral infections such as hepatitis. The present sequence is
CC   the human 9C1LC light chain protein (variable region and J-region)
CC   encoding DNA. This sequence is used in the exemplification of the
CC   invention.
XX
SQ   Sequence 327 BP; 84 A; 89 C; 76 G; 78 T; 0 U; 0 Other;

  Query Match            74.3%;  Score 238.6;  DB 13;  Length 327;
  Best Local Similarity  84.5%;  Pred. No. 7.5e-67;
  Matches  268;  Conservative    0;  Mismatches   49;  Indels    0;  Gaps    0;

Qy          5 TCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64
              ||||| |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          5 TCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64

Qy         65 CTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGAC 124
              |||||||||||||||| ||||||||||   ||||   ||||||||||||||||||||
Db         65 CTTGCCGGGCAAGTCAGGGCATTAGCAGTGCTTTAGCCTGGTATCAGCAGAAACCAGGGA 124

Qy        125 AGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGAT 184
              |  |||||||||| || || ||    || ||       ||||   |||||||     | |
Db        125 AAGCTCCTAAGCTCCTGATCTATGATGCCTCCTATTTGGAAAGTGGGGTCCCATCAAGGT 184

Qy        185 TCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244
              |||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||||||
Db        185 TCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244

Qy        245 ATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGA 304
              ||||||| ||||| |  |||||||||| | | ||||  ||||||||||||||||||||||
Db        245 ATTTTGCAACTTATTACTGTCAACAGTTTAAAAGTTACCCGATCACCTTCGGCCAAGGGA 304

Qy        305 CACGACTGGACATTCAA 321
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Db        305 CACGACTGGAGATTAAA 321


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GenCore version 5.1.6

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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

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Total number of hits satisfying chosen parameters: 68479088

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Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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7: gb_est6:*
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9: gb_gss2:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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4	236.2	73.6	748	6	CB957070	CB957070 AGENCOURT
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7	233	72.6	781	6	CB985512	CB985512 AGENCOURT
8	231.6	72.1	553	6	CD705041	CD705041 EST21568
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10	231.4	72.1	422	2	AW407904	AW407904 UI-HF-BL0
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20	228.2	71.1	708	6	CB956923	CB956923 AGENCOURT
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27	226.6	70.6	754	6	CB986767	CB986767 AGENCOURT
28	226.6	70.6	785	6	CB955817	CB955817 AGENCOURT
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30	226.6	70.6	923	5	BQ882857	BQ882857 AGENCOURT
31	226.4	70.5	740	6	CB987627	CB987627 AGENCOURT
32	226.4	70.5	756	6	CB984720	CB984720 AGENCOURT
33	225.8	70.3	608	2	AW404714	AW404714 UI-HF-BL0
34	225.8	70.3	921	4	BG341239	BG341239 602463904
35	225	70.1	391	2	AW404992	AW404992 UI-HF-BL0
36	225	70.1	460	2	AW405906	AW405906 UI-HF-BL0
37	225	70.1	510	6	CD694557	CD694557 EST11080
38	225	70.1	550	6	CD709957	CD709957 EST26484
39	225	70.1	566	2	AW406081	AW406081 UI-HF-BL0
40	225	70.1	695	6	CD683876	CD683876 EST396 hu
41	225	70.1	724	6	CB959008	CB959008 AGENCOURT
42	225	70.1	742	6	CB984723	CB984723 AGENCOURT
43	225	70.1	759	6	CB986444	CB986444 AGENCOURT
44	225	70.1	785	6	CB986976	CB986976 AGENCOURT
45	225	70.1	820	6	CB956240	CB956240 AGENCOURT

ALIGNMENTS

RESULT 1
AW404894
LOCUS AW404894 353 bp mRNA linear EST 16-FEB-2000
DEFINITION UI-HF-BL0-acn-f-07-0-UI.r1 NIH_MGC_37 Homo sapiens cDNA clone IMAGE:3059844 5', mRNA sequence.
ACCESSION AW404894
VERSION AW404894.1 GI:6923951
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 353)
AUTHORS NIH-MGC http://mgc.nci.nih.gov/.
TITLE National Institutes of Health, Mammalian Gene Collection (MGC)
JOURNAL Unpublished (1999)
COMMENT Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Eco RI site shown at the beginning of the sequence.
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: M.B. Soares Lab
cDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
Seq primer: M13 Forward.
FEATURES Location/Qualifiers
source 1. .353
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:3059844"
/tissue_type="lymph"
/cell_type="germinal center B cells"
/cell_line="MGC85"
/lab_host="DH10B (LTI)"
/clone_lib="NIH_MGC_37"
/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI; Constructed from size fractionated cytoplasmic mRNA (1.5-2.5kb). Directionally cloned. Cells provided by Louis M. Staudt, Ph.D. Library preparation by Maria de Fatima Bonaldo, Ph.D. and M. Bento Soares, Ph.D."
ORIGIN

Query Match 74.3%; Score 238.4; DB 2; Length 353;
Best Local Similarity 85.3%; Pred. No. 5.8e-66;

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  Matches  266;  Conservative    0;  Mismatches   46;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||
Db         42 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTGGGAGACAGAGTCACC 101

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||||||||||||||||||||||||| | ||||||||||||||||||||||||
Db        102 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGTTCTTTAAATTGGTATCAGCAGAAACCA 161

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||   |  | |||| ||| || || ||     ||||| |     | ||   ||||||||
Db        162 GGGAAAGCCCCTAGGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 221

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| |||
Db        222 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 281

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||.|||| |||||||  |||||||||  | ||||||  |||||||||||||||||||
Db        282 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTTCGCCGATCACCTTCGGCCAA 341

Qy        301 GGGACACGACTG 312
              ||||||:||||
Db        342 GGGACACGACTG 353


RESULT 2
CB955875
LOCUS       CB955875                794 bp    mRNA    linear   EST 29-APR-2003
DEFINITION  AGENCOURT_13779371 NIH_MGC_184 Homo sapiens cDNA clone
            IMAGE:30349554 5', mRNA sequence.
ACCESSION   CB955875
VERSION     CB955875.1  GI:30211993
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 794)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Dr. Michael Brownstein and Dr. Miklos Palkovits
             cDNA Library Preparation: CLONTECH Laboratories, Inc.
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Agencourt Bioscience Corporation
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: NDCM144  row: a  column: 19
            High quality sequence stop: 520.
FEATURES             Location/Qualifiers
     source          1. .794
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:30349554"
                     /lab_host="DH10B (T1 phage-resistant)"
                     /clone_lib="NIH_MGC_184"
                     /note="Organ: Pooled-Glandular; Vector: pDNR-LIB; Site_1:
                     SfiI (ggccattatggcc); Site_2: SfiI (ggccgcctcggcc);
                     Library is oligo-dT primed and directionally cloned. cDNA
                     was prepared from a glandular pool of tissues from thyoid,
                     parathyroid, adrenal, cortex and pineal gland.  5' and 3'
                     adaptors were used in cloning as follows: 5' adaptor
                     sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence:
                     5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3' (where B = A,
                     C, or G and N = A, C, G, or T). Average insert size 1.38
                     kb (range 0.60-3.5 kb). 15/15 colonies contained inserts
                     by PCR. This library was enriched for full-length clones
                     and was constructed by Clontech Laboratories (Palo Alto,
                     CA). Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             74.1%;  Score 237.8;  DB 6;  Length 794;
  Best Local Similarity   83.8%;  Pred. No. 1.2e-65;
  Matches  269;  Conservative    0;  Mismatches   52;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         94 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 153

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||  |||||||| ||||||||||||||||||||||||||||
Db        154 ATCACTTGCCGGGCAAGTCAGTACATTAGCACCTATTTAAATTGGTATCAGCAGAAACCA 213

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||   |  | |||||  | || |||||      || | ||    |||   ||||||||
Db        214 GGGAAAGCCCCTAAATTACTGATTTATACTACAACCACTTTAGAACAGGGGGTCCCATCA 273

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||  ||||||| |||||||||| ||| |||||||||||||| | ||||| | |
Db        274 AGGTTCACTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCCGTCTGCAACTT 333

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||||||| | ||||||||||||||  | |||||||  ||| ||||||||||||||
Db        334 GAAGATTTTGCAAATTACTTTTGTCAACAGAGTTACAGTTTCCCGGTCACCTTCGGCCAA 393

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||||||||||| ||| ||
Db        394 GGGACACGACTGGAGATTAAA 414


RESULT 3
BM171889
LOCUS       BM171889                621 bp    mRNA    linear   EST 04-DEC-2001
DEFINITION  imageqc_3_2001/smm27bdff41.x1 NIH_MGC_77 Homo sapiens cDNA clone
            IMAGE:4690877 5', mRNA sequence.
ACCESSION   BM171889
VERSION     BM171889.1  GI:17311452
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 621)
  AUTHORS   Kale,P.I., Harsch,T.J., Folta,P.A., Nelson,D.O., Sanders,C.G. and
            Prange,C.K.
  TITLE     The I.M.A.G.E. Consortium quality control effort: clone
            resequencing for verification
  JOURNAL   Unpublished (2001)
COMMENT     Other_ESTs: BG536784
            Contact: Prange CK
            The I.M.A.G.E. Consortium
            Lawrence Livermore National Laboratory
            Livermore, CA, USA
            Email: help@image.llnl.gov
            This read has been verified (found to hit its original self in the
            correct orientation), as part of the I.M.A.G.E. Consortium quality
            control effort. High quality sequence is defined as having 100 or
            more base pairs with a phred quality value of 20 or greater, where
            a sliding window of 4 base pairs with a phred quality value of 15
            or greater marks the beginning and end of the sequence. For
            information on obtaining this clone, please contact
            info@image.llnl.gov.
            Plate: LLCM1509  row: c  column: 6
            Seq primer: -21m13
            High quality sequence stop: 621.
FEATURES             Location/Qualifiers
     source          1. .621
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
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/clone="IMAGE:4690877"
/lab_host="DH10B (T1 phage-resistant)"
/clone_lib="NIH_MGC_77"
/note="Organ: lung; Vector: pDNR-LIB (Clontech); Site_1: SfiI (ggccgcctcggcc); Site_2: SfiI (ggccattatggcc); 5' and 3' adaptors were used in cloning as follows: 5' adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence: 5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3' (where B = A, C, or G and N = A, C, G, or T). Average insert size 1.9 kb (range 0.5-4.0 kb). 12/15 colonies contained inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo Alto, CA). Note: this is a NIH_MGC Library."

ORIGIN

Query Match 73.6%; Score 236.2; DB 4; Length 621;
Best Local Similarity 83.5%; Pred. No. 3.7e-65;
Matches 268; Conservative 0; Mismatches 53; Indels 0; Gaps 0;

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Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||   |||||||||||||||||||||||||||||| ||||||| ||||||||||||||||
Db         90 GATTGCCAGATGACCCAGTCTCCATCCTCCCTGTCTACATCTGTGGGAGACAGAGTCACC 149

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
                |||||||||||||||||||| ||||| ||  |||||||||||||||||  || ||||||
Db        150 GCCACTTGCCGGGCAAGTCAGAACATTAACAAATATTTAAATTGGTATCAACAAAAACCA 209

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              || ||  | |||||||| || |||||  | ||||| ||   | ||    ||||||||
Db        210 GGGCAAGCCCCTAAGCTCCTGATTTATGGTGCATCCACTTTGCAAAGTGGGGTCCCATCA 269

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | |||||  |||||||  |||||||||| ||| ||||||||||||||||||  ||||| |||
Db        270 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 329

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||||||| |||||||  |||||||| ||  || | | ||||||||||||||||||
Db        330 GAAGATTTTGCAACTTACTACTGTCAACAGACTTTCATTATCCCGATCACCTTCGGCCAA 389

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||| ||| ||
Db        390 GGGACACGACTGGAGATTAAA 410
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RESULT 4
CB957070
LOCUS CB957070 748 bp mRNA linear EST 29-APR-2003
DEFINITION AGENCOURT_13777897 NIH_MGC_184 Homo sapiens cDNA clone IMAGE:30350034 5', mRNA sequence.
ACCESSION CB957070
VERSION CB957070.1 GI:30213187
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 748)
AUTHORS NIH-MGC http://mgc.nci.nih.gov/.
TITLE National Institutes of Health, Mammalian Gene Collection (MGC)
JOURNAL Unpublished (1999)
COMMENT Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Michael Brownstein and Dr. Miklos Palkovits
cDNA Library Preparation: CLONTECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: NDCM145 row: e column: 19
High quality sequence stop: 507.
FEATURES Location/Qualifiers
source 1. .748

/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:30350034"
/lab_host="DH10B (T1 phage-resistant)"
/clone_lib="NIH_MGC_184"
/note="Organ: Pooled-Glandular; Vector: pDNR-LIB; Site_1: SfiI (ggccattatggcc); Site_2: SfiI (ggccgcctcggcc); Library is oligo-dT primed and directionally cloned. cDNA was prepared from a glandular pool of tissues from thyoid, parathyroid, adrenal, cortex and pineal gland. 5' and 3' adaptors were used in cloning as follows: 5' adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence: 5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3' (where B = A, C, or G and N = A, C, G, or T). Average insert size 1.38 kb (range 0.60-3.5 kb). 15/15 colonies contained inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo Alto, CA). Note: this is a NIH_MGC Library."

ORIGIN

Query Match 73.6%; Score 236.2; DB 6; Length 748;
Best Local Similarity 83.5%; Pred. No. 4e-65;
Matches 268; Conservative 0; Mismatches 53; Indels 0; Gaps 0;

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Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  |||||||||||||||||||||||||||||||||||||||| | ||||||||||||||||
Db         89 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTATTGGAGACAGAGTCACC 148

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||||||||||||||||||||||| ||| ||||||||||||||||||||||||||||
Db        149 ATCACTTGCCGGGCAAGTCAGAGCATTACCAGTTATTTAAATTGGTATCAGCAGAAACCA 208

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  | ||  ||||| || || || ||    |||||       | ||   ||||||||
Db        209 GGGGAACCCCCTAACCTCCTGATCTATGCTGCATCCCGTTTGCAAAGTGGGGTCCCATCA 268

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| ||||||| ||||| |||| ||| |||||||||||||||||| ||||| |||
Db        269 AGGTTCAGTGGCAGTGGATCTGAGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 328

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              || |||||||| |||||||| |||||||||  | ||| |   ||||||||||||||||||
Db        329 GATGATTTTGCAACTTACTTCTGTCAACAGAGTTACAATACCCCGATCACCTTCGGCCAA 388

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||| ||| ||
Db        389 GGGACACGACTGGAGATTAAA 409
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RESULT 5
BG742662
LOCUS BG742662 826 bp mRNA linear EST 15-MAY-2001
DEFINITION 602633264F1 NCI_CGAP_Skn3 Homo sapiens cDNA clone IMAGE:4778726 5', mRNA sequence.
ACCESSION BG742662
VERSION BG742662.1 GI:14053315
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 826)
AUTHORS NIH-MGC http://mgc.nci.nih.gov/.
TITLE National Institutes of Health, Mammalian Gene Collection (MGC)
JOURNAL Unpublished (1999)
COMMENT Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: James Cleaver, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be

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            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLAM10634  row: o  column: 15
            High quality sequence stop: 824.
FEATURES              Location/Qualifiers
     source           1. .826
                      /organism="Homo sapiens"
                      /mol_type="mRNA"
                      /db_xref="taxon:9606"
                      /clone="IMAGE:4778726"
                      /lab_host="DH10B (T1 phage-resistant)"
                      /clone_lib="NCI_CGAP_Skn3"
                      /note="Organ: skin; Vector: pCMV-SPORT6; Site_1: NotI;
                      Site_2: SalI; Cloned unidirectionally.  Primer: Oligo dT.
                      Average insert size 1.5kb.  Library constructed by Life
                      Technologies. Note: this is a NCI_CGAP Library."
ORIGIN

  Query Match             73.1%;  Score 234.6;  DB 4;  Length 826;
  Best Local Similarity   83.2%;  Pred. No. 1.4e-64;
  Matches  267;  Conservative    0;  Mismatches   54;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         55 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 114

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||||||||||||||||||||||| |||||||||||||| |||||||||| ||
Db        115 ATCACTTGCCGGGCAAGTCAGAGCATTAGCACCTATTTAAATTGGTTTCAGCAGAAATCA 174

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||| || || ||  |  | ||||| |  | | |    ||||||||
Db        175 GGGAAAGCCCCTAAACTCCTGATCCATGGTGCATCCAGTCTGCAGAGTGGGGTCCCATCA 234

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | |||||| ||||||| |||||||||| ||| |||||||||||||| || ||||| |||
Db        235 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGTAGTCTGCAACCT 294

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| | |||||  |||||||||  | |||||  | |||| ||||||||||||
Db        295 GAAGATTTTGCAATTTACTACTGTCAACAGGGTTACAGTCAGGCGATTACCTTCGGCCAA 354

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||| || ||| ||
Db        355 GGGACACGACTAGAGATTAAA 375


RESULT 6
CD690167
LOCUS       CD690167                 709 bp    mRNA    linear   EST 25-JUN-2003
DEFINITION  EST6690 human nasopharynx Homo sapiens cDNA, mRNA sequence.
ACCESSION   CD690167
VERSION     CD690167.1  GI:32210659
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 709)
  AUTHORS   Liu,X.-Q., Zhou,Y., Zhang,L.-J., Xu,H., Chen,H.-K., Pan,Z.-G. and
            Zeng,Y.-X.
  TITLE     Transcriptional Gene Expression Profile of Human Nasopharynx
  JOURNAL   Unpublished (2003)
COMMENT     Contact: YiXin Zeng
            Cancer Center
            Sun Yat-sen University
            651 DongFeng Road East, GuangZhou 510060, China
            Tel: 86-1380-9770-743
            Fax: 86-20-8775-4506
            Email: yxzeng@gzsums.edu.cn.
FEATURES              Location/Qualifiers
     source           1. .709
                      /organism="Homo sapiens"
                      /mol_type="mRNA"
                      /db_xref="taxon:9606"
                      /tissue_type="normal nasopharynx"
                      /clone_lib="human nasopharynx"
                      /note="ESTs generated from a normal nasopharynx cDNA
                      library from southern Chinese"
ORIGIN

  Query Match             72.6%;  Score 233;  DB 6;  Length 709;
  Best Local Similarity   82.9%;  Pred. No. 4.2e-64;
  Matches  266;  Conservative    0;  Mismatches   55;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        123 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTGGGAGACAGAGTCACC 182

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||| ||| ||||||  |||||||| ||||||||||||||||||||||||||||
Db        183 ATCACTTGCCAGGCGAGTCAGGACATTAGCAACTATTTAAATTGGTATCAGCAGAAACCG 242

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||||| || || |||   |||| |    ||||   ||||||||
Db        243 GGGAAAGCCCCTAAGCTCCTGATCTACGATGCATCCAATTTGGAAATAGGGGTCCCATCA 302

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | |||| || |||| |||||||||| |||  ||| |||||||||||||||||||||||
Db        303 AGGTTCAGTGGAAGTGGATCTGGGACAGATTTTACTTTCACCATCAGCAGCCTGCAGCCT 362

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||| |||| || || |  |||||||||| ||| | | | | ||||||||||||||||
Db        363 GAAGATATTGCAACGTATTACTGTCAACAGTATGATAATCTCCGGATCACCTTCGGCCAA 422

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||| ||| ||
Db        423 GGGACACGACTGGAGATTAAA 443


RESULT 7
CB985512
LOCUS       CB985512                 781 bp    mRNA    linear   EST 01-MAY-2003
DEFINITION  AGENCOURT_13672145 NIH_MGC_184 Homo sapiens cDNA clone
            IMAGE:30327573 5', mRNA sequence.
ACCESSION   CB985512
VERSION     CB985512.1  GI:30280036
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 781)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Dr. Michael Brownstein and Dr. Miklos Palkovits
             cDNA Library Preparation: CLONTECH Laboratories, Inc.
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Agencourt Bioscience Corporation
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: NDCM136  row: m  column: 22
            High quality sequence stop: 390.
FEATURES              Location/Qualifiers
     source           1. .781
                      /organism="Homo sapiens"
                      /mol_type="mRNA"
                      /db_xref="taxon:9606"
                      /clone="IMAGE:30327573"
                      /lab_host="DH10B (T1 phage-resistant)"
                      /clone_lib="NIH_MGC_184"
                      /note="Organ: Pooled-Glandular; Vector: pDNR-LIB; Site_1:
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SfiI (ggccattatggcc); Site_2: SfiI (ggccgcctcggcc); Library is oligo-dT primed and directionally cloned. cDNA was prepared from a glandular pool of tissues from thyoid, parathyroid, adrenal, cortex and pineal gland. 5' and 3' adaptors were used in cloning as follows: 5' adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence: 5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3' (where B = A, C, or G and N = A, C, G, or T). Average insert size 1.38 kb (range 0.60-3.5 kb). 15/15 colonies contained inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo Alto, CA). Note: this is a NIH_MGC Library."

ORIGIN

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  Query Match            72.6%;  Score 233;  DB 6;  Length 781;
  Best Local Similarity  82.9%;  Pred. No. 4.4e-64;
  Matches  266;  Conservative    0;  Mismatches   55;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         89 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 148

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||  ||||||||||||||  || ||||  ||||||||||||||||||  |||||  |||
Db        149 ATCACTTGTCGGGCAAGTCAGAGAATAAGCACCTATTTAAATTGGTATCAACAGAAGCCA 208

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||  || || || ||    |||||  |    | ||   ||||||||
Db        209 GGGAAAGCCCCTAACCTCCTAATCTATACTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 268

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | |||||  |||||||  ||||||||||  |||  ||||||||||||||||||  |||||  |||
Db        269 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 328

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||||||  |||||  |  |||||||||  |  |||  |    |||  ||||||||||||||
Db        329 GAAGATTTTGCCACTTATTACTGTCAACAGAGTTACAATACCCCGGTCACCTTCGGCCAA 388

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||||||  ||
Db        389 GGGACACGACTGGACATTAAA 409
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RESULT 8
CD705041
LOCUS CD705041 553 bp mRNA linear EST 25-JUN-2003
DEFINITION EST21568 human nasopharynx Homo sapiens cDNA, mRNA sequence.
ACCESSION CD705041
VERSION CD705041.1 GI:32235671
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 553)
AUTHORS Liu,X.-Q., Zhou,Y., Zhang,L.-J., Xu,H., Chen,H.-K., Pan,Z.-G. and Zeng,Y.-X.
TITLE Transcriptional Gene Expression Profile of Human Nasopharynx
JOURNAL Unpublished (2003)
COMMENT Contact: YiXin Zeng
Cancer Center
Sun Yat-sen University
651 DongFeng Road East, GuangZhou 510060, China
Tel: 86-1380-9770-743
Fax: 86-20-8775-4506
Email: yxzeng@gzsums.edu.cn.
FEATURES Location/Qualifiers
source 1. .553
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/tissue_type="normal nasopharynx"
/clone_lib="human nasopharynx"
/note="ESTs generated from a normal nasopharynx cDNA library from southern Chinese"

ORIGIN

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  Query Match            72.1%;  Score 231.6;  DB 6;  Length 553;
  Best Local Similarity  83.0%;  Pred. No. 1.1e-63;
  Matches  264;  Conservative    0;  Mismatches   54;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        123 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCTTCTGTAGGAGACAGAGTCACC 182

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||  ||||||||||||||||||||||||||
Db        183 ATCACTTGCCGGGCAAGTCAGAGCATTAGGTTCTATTTAAATTGGTATCAGCAGAAACCA 242

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||  || || || ||    |||||  |    | ||   ||||||||
Db        243 GGGAAAGCCCCTAATCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATTA 302

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | | |||  |||||||  ||||||||||  |||  ||||||||||||||||  |||  |||||  |||
Db        303 AGGTACAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCACCAGTCTGCAACCT 362

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||||||  |||||||  |||||||||  ||  ||  |||  ||  ||||  |||||||||
Db        363 GAAGATTTTGCAACTTACTACTGTCAACAGACTTACGGTTCCCCTCTCACTTTCGGCCAA 422

Qy        301 GGGACACGACTGGACATT 318
              ||||||||||||||  |||
Db        423 GGGACACGACTGGAGATT 440
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RESULT 9
AW406227
LOCUS AW406227 421 bp mRNA linear EST 16-FEB-2000
DEFINITION UI-HF-BL0-acp-c-08-0-UI.r1 NIH_MGC_37 Homo sapiens cDNA clone IMAGE:3059726 5', mRNA sequence.
ACCESSION AW406227
VERSION AW406227.1 GI:6925284
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 421)
AUTHORS NIH-MGC http://mgc.nci.nih.gov/.
TITLE National Institutes of Health, Mammalian Gene Collection (MGC)
JOURNAL Unpublished (1999)
COMMENT Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Eco RI site shown at the beginning of the sequence.
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: M.B. Soares Lab
cDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Seq primer: M13 Forward.
FEATURES Location/Qualifiers
source 1. .421
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:3059726"
/tissue_type="lymph"
/cell_type="germinal center B cells"
/cell_line="MGC85"
/lab_host="DH10B (LTI)"
/clone_lib="NIH_MGC_37"
/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI; Constructed from size fractionated cytoplasmic mRNA

(1.5-2.5kb). Directionally cloned. Cells provided by Louis M. Staudt, Ph.D. Library preparation by Maria de Fatima Bonaldo, Ph.D. and M. Bento Soares, Ph.D."

ORIGIN

Query Match 72.1%; Score 231.4; DB 2; Length 421;
Best Local Similarity 82.6%; Pred. No. 1.2e-63;
Matches 265; Conservative 0; Mismatches 56; Indels 0; Gaps 0;

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Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||| ||| ||||||||||||||||||||| |||||||
Db         10 GACATCCAGATGACCCAGTCTCCATCTTCCGTGTCTGCATCTGTAGGAGACACAGTCACC 69

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||||| |||||| |  || ||||||  |||   ||||||||||||||||||
Db         70 ATCACTTGTCGGGCGAGTCAGGGTCTTGGCAGCTGGTTAGCCTGGTATCAGCAGAAACCA 129

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||     | |||||||| ||  | ||    ||||| |    | ||   ||||||||
Db        130 GGGACTGCCCCTAAGCTCCTGCTCTATGCTGCATCCAATTTGCAAAGTGGGGTCCCATCA 189

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | |||||||||||| |||||||||| ||| ||||||||||||  ||||||||||||
Db        190 AGGTTCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCGACAGCCTGCAGCCT 249

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||||||||||| |||||||||| || ||| ||| ||||||||||||||||||
Db        250 GAAGATTTTGCTACTTACTATTGTCAACAGGCTAACACTTTCCCGATCACCTTCGGCCAA 309

Qy        301 GGGACACGACTGGACATTCAA 321
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Db        310 GGGACACGACTGGACATTAAA 330
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RESULT 10
AW407904
LOCUS AW407904 422 bp mRNA linear EST 16-FEB-2000
DEFINITION UI-HF-BL0-add-a-01-0-UI.r2 NIH_MGC_37 Homo sapiens cDNA clone IMAGE:3061128 5', mRNA sequence.
ACCESSION AW407904
VERSION AW407904.1 GI:6926961
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 422)
AUTHORS NIH-MGC http://mgc.nci.nih.gov/.
TITLE National Institutes of Health, Mammalian Gene Collection (MGC)
JOURNAL Unpublished (1999)
COMMENT Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Eco RI site shown at the beginning of the sequence.
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: M.B. Soares Lab
cDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Seq primer: M13 Forward.
FEATURES Location/Qualifiers
source 1. .422
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:3061128"
/tissue_type="lymph"
/cell_type="germinal center B cells"
/cell_line="MGC85"
/lab_host="DH10B (LTI)"
/clone_lib="NIH_MGC_37"
/note="Vector: pT7T3-Pac; Site_1: NotI; Site_2: Eco RI;

Constructed from size fractionated cytoplasmic mRNA (1.5-2.5kb). Directionally cloned. Cells provided by Louis M. Staudt, Ph.D. Library preparation by Maria de Fatima Bonaldo, Ph.D. and M. Bento Soares, Ph.D."

ORIGIN

Query Match 72.1%; Score 231.4; DB 2; Length 422;
Best Local Similarity 82.6%; Pred. No. 1.2e-63;
Matches 265; Conservative 0; Mismatches 56; Indels 0; Gaps 0;

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Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         23 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 82

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         83 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 142

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||||||  || ||  ||    ||||| |    | ||   ||||||||
Db        143 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 202

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | |||||  |||||||  ||||||||||  ||| ||||||||||||||||||  ||||| |||
Db        203 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 262

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||||||| |||||||  |||||||||  | |||||    |||   || |||||||||
Db        263 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGTGGACGTTCGGCCAA 322

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||     |||| ||  ||
Db        323 GGGACCAAGGTGGAAATCAAA 343
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RESULT 11
CD707755
LOCUS CD707755 521 bp mRNA linear EST 25-JUN-2003
DEFINITION EST24282 human nasopharynx Homo sapiens cDNA, mRNA sequence.
ACCESSION CD707755
VERSION CD707755.1 GI:32238385
KEYWORDS EST.
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 521)
AUTHORS Liu,X.-Q., Zhou,Y., Zhang,L.-J., Xu,H., Chen,H.-K., Pan,Z.-G. and Zeng,Y.-X.
TITLE Transcriptional Gene Expression Profile of Human Nasopharynx
JOURNAL Unpublished (2003)
COMMENT Contact: YiXin Zeng
Cancer Center
Sun Yat-sen University
651 DongFeng Road East, GuangZhou 510060, China
Tel: 86-1380-9770-743
Fax: 86-20-8775-4506
Email: yxzeng@gzsums.edu.cn.
FEATURES Location/Qualifiers
source 1. .521
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/tissue_type="normal nasopharynx"
/clone_lib="human nasopharynx"
/note="ESTs generated from a normal nasopharynx cDNA library from southern Chinese"

ORIGIN

Query Match 72.1%; Score 231.4; DB 6; Length 521;
Best Local Similarity 82.6%; Pred. No. 1.3e-63;
Matches 265; Conservative 0; Mismatches 56; Indels 0; Gaps 0;

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Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         91 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 150

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||| ||| ||||||  ||| | |  ||||||||||||||||||||||||||||||
Db        151 ATCACTTGCCAGGCGAGTCAGGACATAAACTACTATTTAAATTGGTATCAGCAGAAACCA 210

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||||||  || || |||   ||||| |    |||| | ||||||  |
Db        211 GGGAAAGCCCCTAAGCTCCTGATCTACGATGCATCCATTTTGGAAACAGGGGTCTCATCA 270

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| || |||| |||||||||| |||  ||| |||||||||| ||||||||||||
Db        271 AGGTTCAGTGGAAGTGGATCTGGGACAGATTTTACTTTCACCATCACCAGCCTGCAGCCT 330

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||| |||| || || |  |||||||||| ||| | | | ||||||||||||||||||
Db        331 GAAGATATTGCAACATATTACTGTCAACAGTTTGATAATCTCCCGATCACCTTCGGCCAA 390

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||| ||| ||
Db        391 GGGACACGACTGGAGATTAAA 411


RESULT 12
BG756289
LOCUS       BG756289                 587 bp    mRNA    linear   EST 15-MAY-2001
DEFINITION  602713607F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4853953 5',
            mRNA sequence.
ACCESSION   BG756289
VERSION     BG756289.1  GI:14066942
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 587)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
             cDNA Library Preparation: Ling Hong/Rubin Laboratory
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1700  row: n  column: 02
            High quality sequence stop: 586.
FEATURES             Location/Qualifiers
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                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:4853953"
                     /tissue_type="primary B-cells from tonsils (cell line)"
                     /lab_host="DH10B (phage-resistant)"
                     /clone_lib="NIH_MGC_48"
                     /note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
                     Site_2: EcoRI; cDNA made by oligo-dT priming.
                     Directionally cloned into EcoRI/XhoI sites using the
                     following 5' adaptor: GGCACGAG(G). Size-selected >500bp
                     for average insert size 1.8kb. Library constructed by Ling
                     Hong in the laboratory of Gerald M. Rubin (University of
                     California, Berkeley) using  ZAP-cDNA synthesis kit
                     (Stratagene) and Superscript II RT (Life Technologies).
                     Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             72.1%;  Score 231.4;  DB 4;  Length 587;
  Best Local Similarity   82.6%;  Pred. No. 1.3e-63;
  Matches  265;  Conservative    0;  Mismatches   56;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         77 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTGGGAGACAGAGTCACC 136

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||| ||| ||||||   |||||   |||||||||||||||||||||||||||||
Db        137 ATCACTTGCCAGGCGAGTCAGGATATTAGTTACTATTTAAATTGGTATCAGCAGAAACCA 196

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||| |  | ||||||||  || || |||   ||||| |    ||||   ||||||||
Db        197 GGAAAAGCCCCTAAGCTCCTGATCTACGATGCATCCAATTTGGAAATAGGGGTCCCATCA 256

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| ||  ||| |||||||||| |||  ||| ||||||||||||||||||||||||
Db        257 AGGTTCAGTGGGCGTGGATCTGGGACAGATTTTACTTTCACCATCAGCAGCCTGCAGCCT 316

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||| |||| || || | ||||||||  |  ||| ||| | ||||||||||||||||||
Db        317 GAAGATATTGCAACATATTATTGTCAACGGCATGATAGTCTCCCGATCACCTTCGGCCAA 376

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||| ||| ||
Db        377 GGGACACGACTGGAAATTAAA 397


RESULT 13
CD690145
LOCUS       CD690145                 624 bp    mRNA    linear   EST 25-JUN-2003
DEFINITION  EST6668 human nasopharynx Homo sapiens cDNA, mRNA sequence.
ACCESSION   CD690145
VERSION     CD690145.1  GI:32210615
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 624)
  AUTHORS   Liu,X.-Q., Zhou,Y., Zhang,L.-J., Xu,H., Chen,H.-K., Pan,Z.-G. and
            Zeng,Y.-X.
  TITLE     Transcriptional Gene Expression Profile of Human Nasopharynx
  JOURNAL   Unpublished (2003)
COMMENT     Contact: YiXin Zeng
            Cancer Center
            Sun Yat-sen University
            651 DongFeng Road East, GuangZhou 510060, China
            Tel: 86-1380-9770-743
            Fax: 86-20-8775-4506
            Email: yxzeng@gzsums.edu.cn.
FEATURES             Location/Qualifiers
     source          1. .624
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /tissue_type="normal nasopharynx"
                     /clone_lib="human nasopharynx"
                     /note="ESTs generated from a normal nasopharynx cDNA
                     library from southern Chinese"
ORIGIN

  Query Match             72.1%;  Score 231.4;  DB 6;  Length 624;
  Best Local Similarity   82.6%;  Pred. No. 1.3e-63;
  Matches  265;  Conservative    0;  Mismatches   56;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        114 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 173

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        174 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 233
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Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
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Db        234 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 293

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Db        294 AGATTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 353

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  ||||||||  | |||||   || |  || || |||||
Db        354 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCTAGAACTTTTGGCCAG 413

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||    |||||| ||  ||
Db        414 GGGACCAAGCTGGAGATCAAA 434


RESULT 14
BG536784
LOCUS       BG536784                 570 bp    mRNA    linear   EST 03-APR-2001
DEFINITION  602566318F1 NIH_MGC_77 Homo sapiens cDNA clone IMAGE:4690877 5',
            mRNA sequence.
ACCESSION   BG536784
VERSION     BG536784.1  GI:13528330
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 570)
  AUTHORS   NIH-MGC http://mgc.nci.nih.gov/.
  TITLE     National Institutes of Health, Mammalian Gene Collection (MGC)
  JOURNAL   Unpublished (1999)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: CLONTECH Laboratories, Inc.
             cDNA Library Preparation: CLONTECH Laboratories, Inc.
             cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
             DNA Sequencing by: Incyte Genomics, Inc.
             Clone distribution: MGC clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            http://image.llnl.gov
            Plate: LLCM1509  row: c  column: 06
            High quality sequence stop: 568.
FEATURES             Location/Qualifiers
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                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /clone="IMAGE:4690877"
                     /lab_host="DH10B (T1 phage-resistant)"
                     /clone_lib="NIH_MGC_77"
                     /note="Organ: lung; Vector: pDNR-LIB (Clontech); Site_1:
                     SfiI (ggccgcctcggcc); Site_2: SfiI (ggccattatggcc); 5' and
                     3' adaptors were used in cloning as follows: 5' adaptor
                     sequence:  5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence:
                     5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3' (where B = A,
                     C, or G and N = A, C,  G, or T).  Average insert size 1.9
                     kb (range 0.5-4.0 kb). 12/15 colonies contained inserts
                     by PCR. This library was enriched for full-length clones
                     and was constructed by Clontech  Laboratories (Palo Alto,
                     CA). Note: this is a NIH_MGC Library."
ORIGIN

  Query Match             71.8%;  Score 230.6;  DB 4;  Length 570;
  Best Local Similarity   84.4%;  Pred. No. 2.4e-63;
  Matches  271;  Conservative    0;  Mismatches   49;  Indels    1;  Gaps    1;

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Db         91 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTACATCTGTGGGAGACAGAGTCACC 150

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||| ||||| ||  |||||||||||||||| || ||||||
Db        151 GTCACTTGCCGGGCAAGTCAGAACATTAACAAATATTTAAATTGGTATCAACAAAAACCA 210

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |||| |||||||| || |||||  | |||||  ||   | ||   ||||||||
Db        211 GGCAAGCC-CCTAAGCTCCTGATTTATGGTGCATCCACTTTGCAAAGTGGGGTCCCATCA 269

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| ||||||| |||||||||| ||| ||||||||||||||||| ||||| |||
Db        270 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 329

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||||  ||  || | | ||||||||||||||||||
Db        330 GAAGATTTTGCAACTTACTACTGTCAACAGACTTTCATTATCCCGATCACCTTCGGCCAA 389

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||||||||||| ||| ||
Db        390 GGGACACGACTGGAGATTAAA 410


RESULT 15
CD683960
LOCUS       CD683960                 486 bp    mRNA    linear   EST 25-JUN-2003
DEFINITION  EST480 human nasopharynx Homo sapiens cDNA, mRNA sequence.
ACCESSION   CD683960
VERSION     CD683960.1  GI:32198500
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 486)
  AUTHORS   Liu,X.-Q., Zhou,Y., Zhang,L.-J., Xu,H., Chen,H.-K., Pan,Z.-G. and
            Zeng,Y.-X.
  TITLE     Transcriptional Gene Expression Profile of Human Nasopharynx
  JOURNAL   Unpublished (2003)
COMMENT     Contact: YiXin Zeng
            Cancer Center
            Sun Yat-sen University
            651 DongFeng Road East, GuangZhou 510060, China
            Tel: 86-1380-9770-743
            Fax: 86-20-8775-4506
            Email: yxzeng@gzsums.edu.cn.
FEATURES             Location/Qualifiers
     source          1. .486
                     /organism="Homo sapiens"
                     /mol_type="mRNA"
                     /db_xref="taxon:9606"
                     /tissue_type="normal nasopharynx"
                     /clone_lib="human nasopharynx"
                     /note="ESTs generated from a normal nasopharynx cDNA
                     library from southern Chinese"
ORIGIN

  Query Match             71.6%;  Score 229.8;  DB 6;  Length 486;
  Best Local Similarity   82.2%;  Pred. No. 4e-63;
  Matches  264;  Conservative    0;  Mismatches   57;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  |||||||||||||| |||||| ||| ||||||||||||||||||||||||||||||
Db        134 GACATCCAGATGACCCAGTTTCCATCTTCCGTGTCTGCATCTGTAGGAGACAGAGTCACC 193

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||||| ||||||   ||||||||||  |||   ||||||||||||||||||
Db        194 ATCACTTGTCGGGCGAGTCAGGATATTAGCAGCTGGTTAGCCTGGTATCAGCAGAAACCA 253

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||| | || || ||     ||||| |       ||   ||||||||
Db        254 GGGAAAGCCCCTAAGGTCCTGATCTATGCTGCATCCAATTTACAAAGTGGGGTCCCATCA 313

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | |||||||||||| |||||||||||  | ||||||||||||||||||||||||||||||
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Db        314 AGGTTCAGCGGCAGTGGATCTGGGACAAAATTCACTCTCACCATCAGCAGCCTGCAGCCT 373

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||||| ||||| | |||||||||||  | |||| || ||||||||||||||||||
Db        374 GAAGATTTTGCAACTTATTATTGTCAACAGGGTAACAGGTTCCCGATCACCTTCGGCCAA 433

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||||||||||| ||| ||
Db        434 GGGACACGACTGGAGATTAAA 454
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Job time : 2584.67 secs

GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: September 11, 2005, 19:40:23 ; Search time 98.7318 Seconds
(without alignments)
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Title: US-09-403-107-147
Perfect score: 321
Sequence: 1 gagctccagatgacccagtc.....ggacacgactggacattcaa 321

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1202784 seqs, 818138359 residues

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:*
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6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	240.8	75.0	321	3	US-09-240-274-106	Sequence 106, App
2	236.2	73.6	321	2	US-08-378-939-13	Sequence 13, Appl
3	231.4	72.1	720	4	US-09-192-854-1	Sequence 1, Appli
4	228.2	71.1	714	4	US-09-472-087-62	Sequence 62, Appl
5	224	69.8	321	3	US-09-240-274-216	Sequence 216, App
6	223.2	69.5	321	3	US-09-240-274-102	Sequence 102, App
7	223.2	69.5	321	3	US-09-240-274-109	Sequence 109, App
8	223.2	69.5	321	3	US-09-240-274-215	Sequence 215, App
9	223.2	69.5	321	3	US-09-240-274-217	Sequence 217, App
10	223.2	69.5	321	3	US-09-240-274-218	Sequence 218, App
11	221.8	69.1	974	4	US-09-859-053-29	Sequence 29, Appl
12	221.6	69.0	321	3	US-09-240-274-199	Sequence 199, App
13	220.2	68.6	388	3	US-09-042-353-358	Sequence 358, App
14	220.2	68.6	388	3	US-08-758-417A-206	Sequence 206, App
15	220	68.5	321	3	US-09-240-274-107	Sequence 107, App
16	218.4	68.0	321	3	US-09-240-274-211	Sequence 211, App
17	218.4	68.0	321	3	US-09-240-274-221	Sequence 221, App
18	218.4	68.0	321	3	US-09-240-274-222	Sequence 222, App
19	217.8	67.9	451	4	US-09-472-087-50	Sequence 50, Appl
20	217	67.6	439	3	US-09-042-353-360	Sequence 360, App
21	217	67.6	439	3	US-08-758-417A-208	Sequence 208, App
22	216.8	67.5	321	3	US-09-240-274-201	Sequence 201, App
23	216.2	67.4	672	4	US-09-456-090A-47	Sequence 47, Appl
24	216.2	67.4	672	4	US-09-453-234-47	Sequence 47, Appl
25	216	67.3	321	3	US-09-240-274-205	Sequence 205, App
26	215.4	67.1	324	2	US-08-378-939-31	Sequence 31, Appl
27	215.4	67.1	324	2	US-08-378-939-33	Sequence 33, Appl
28	215.4	67.1	387	3	US-08-803-085-3	Sequence 3, Appli
29	215.4	67.1	19040	3	US-09-343-485A-3	Sequence 3, Appli
30	215.2	67.0	321	3	US-09-240-274-105	Sequence 105, App
31	215.2	67.0	321	3	US-09-240-274-113	Sequence 113, App
32	215.2	67.0	327	4	US-09-627-896B-25	Sequence 25, Appl
33	213.8	66.6	324	2	US-08-378-939-17	Sequence 17, Appl
34	213.8	66.6	420	3	US-09-042-353-420	Sequence 420, App
35	213.8	66.6	420	3	US-08-758-417A-220	Sequence 220, App
36	213.8	66.6	3819	3	US-09-042-353-393	Sequence 393, App
37	213.8	66.6	3819	3	US-08-758-417A-243	Sequence 243, App
38	213.6	66.5	321	3	US-09-240-274-108	Sequence 108, App
39	213.6	66.5	321	3	US-09-240-274-203	Sequence 203, App
40	213.6	66.5	402	4	US-09-472-087-49	Sequence 49, Appl
41	213.2	66.4	390	2	US-08-646-367-2	Sequence 2, Appli
42	212.2	66.1	321	1	US-08-488-376-12	Sequence 12, Appl
43	212.2	66.1	321	2	US-08-634-223-12	Sequence 12, Appl
44	212.2	66.1	321	2	US-08-634-224-12	Sequence 12, Appl
45	212.2	66.1	321	2	US-08-634-400-12	Sequence 12, Appl

ALIGNMENTS

RESULT 1
US-09-240-274-106
; Sequence 106, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
; APPLICANT: Siegel, Donald L.
; TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
; TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
; FILE REFERENCE: 09596-42U2
; CURRENT APPLICATION NUMBER: US/09/240,274
; CURRENT FILING DATE: 1999-01-29
; EARLIER APPLICATION NUMBER: 60/081,380
; EARLIER FILING DATE: 1998-04-10
; EARLIER APPLICATION NUMBER: 60/028,550
; EARLIER FILING DATE: 1996-10-11
; NUMBER OF SEQ ID NOS: 224
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 106
; LENGTH: 321
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: anti-Rh(D) chain I06
US-09-240-274-106

Query Match 75.0%; Score 240.8; DB 3; Length 321;
Best Local Similarity 85.1%; Pred. No. 9e-75;
Matches 269; Conservative 0; Mismatches 47; Indels 0; Gaps 0;

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Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
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Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
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Db         63 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
                | |||||||| || || ||    ||||| |     | ||    ||||||||      | ||
Db        123 AGCCCCTAAGCTCCTGATCTATGCCGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
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Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| |||||||  |||||||||  | |||||  ||||||||||||||||||||||
Db        243 TTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAAGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
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Db        303 ACGACTGGAGATTAAA 318


RESULT 2
US-08-378-939-13
; Sequence 13, Application US/08378939
; Patent No. 5876961
;  GENERAL INFORMATION:
;    APPLICANT:  CROWE, JAMES SCOTT
;    APPLICANT:  LEWIS, ALAN PETER
;    TITLE OF INVENTION:  PRODUCTION OF ANTIBODIES
;    NUMBER OF SEQUENCES:  46
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  ROTHWELL, FIGG, ERNST & KURZ
;      STREET:  555 THIRTEENTH ST. N.W.
;      CITY:  WASHINGTON
;      STATE:  D. C.
;      COUNTRY:  U.S.
;      ZIP:  20004
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/378,939
;      FILING DATE:
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/952640
;      FILING DATE:  01-DEC-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  ERNST, BARBARA G
;      REGISTRATION NUMBER:  30,377
;      REFERENCE/DOCKET NUMBER:  1808-118
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202) 783-6040
;      TELEFAX:  (202) 783-6031
;  INFORMATION FOR SEQ ID NO:  13:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  321 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  both
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  1..321
US-08-378-939-13

  Query Match             73.6%;  Score 236.2;  DB 2;  Length 321;
  Best Local Similarity   83.5%;  Pred. No. 3.8e-73;
  Matches  268;  Conservative    0;  Mismatches   53;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||  |||||||||||||||| |||||||||
Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAATTATTTAAATTGGTATCAACAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||||| || || ||    ||||| |    | ||   |||||| |
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCACATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| ||||||| |||||||||| | | ||||||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGACTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||| ||| |||||||  |||||||||  | |||||    | ||||||||||||||||
Db        241 GAAGATTCTGCAACTTACTACTGTCAACAGAGTTACAGTACCCTGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 3
US-09-192-854-1
; Sequence 1, Application US/09192854
; Patent No. 6696245
; GENERAL INFORMATION:
;  APPLICANT: Winter, Greg
;  APPLICANT: Tomlinson, Ian
;  TITLE OF INVENTION: Methods for Selecting Functional Peptides
;  FILE REFERENCE: 3789/72916
;  CURRENT APPLICATION NUMBER: US/09/192,854
;  CURRENT FILING DATE: 1998-11-17
;  EARLIER APPLICATION NUMBER: 60/066,729
;  EARLIER FILING DATE: 1997-11-21
;  NUMBER OF SEQ ID NOS: 212
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
;   LENGTH: 720
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-192-854-1

  Query Match             72.1%;  Score 231.4;  DB 4;  Length 720;
  Best Local Similarity   82.6%;  Pred. No. 2.9e-71;
  Matches  265;  Conservative    0;  Mismatches   56;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        397 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 456

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        457 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 516

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||||| || || ||    ||||| |    | ||   |||||| |
Db        517 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 576

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| ||||||| |||||||||| ||| ||||||||||||||||||| ||||| |||
Db        577 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 636

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||||  | |||||   || |  || |||||||||
Db        637 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCTAATACGTTCGGCCAA 696

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||     |||| ||  ||
Db        697 GGGACCAAGGTGGAAATCAAA 717


RESULT 4
US-09-472-087-62
; Sequence 62, Application US/09472087
; Patent No. 6682736
; GENERAL INFORMATION:
;  APPLICANT: HANSON, DOUGLAS C.
;  APPLICANT:  NEVEU, MARK J.
;  APPLICANT:  MUELLER, EILLEN E.
;  APPLICANT:  HANKE, JEFFREY H.
;  APPLICANT:  GILMAN, STEVEN C.
;  APPLICANT:  DAVIS, C. GEOFFREY
;  APPLICANT:  CORVALAN, JOSE R.
;  TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO CTLA-4
;  FILE REFERENCE: ABX-PF1
```

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;  CURRENT APPLICATION NUMBER: US/09/472,087
;  CURRENT FILING DATE:  1999-12-23
;  PRIOR APPLICATION NUMBER: 60/113,647
;  PRIOR FILING DATE: 1998-12-23
;  NUMBER OF SEQ ID NOS: 147
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 62
;   LENGTH: 714
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-472-087-62

  Query Match            71.1%;  Score 228.2;  DB 4;  Length 714;
  Best Local Similarity  81.9%;  Pred. No. 3.9e-70;
  Matches  263;  Conservative    0;  Mismatches   58;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         67 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 126

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||||||||||||||||||||| |||||||||| ||||||||||||||||||||
Db        127 ATCACTTGCCGGGCAAGTCAGAGCATTAACAGCTATTTAGATTGGTATCAGCAGAAACCA 186

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||| || || || ||    ||||| |    | ||   |||||||
Db        187 GGGAAAGCCCCTAAACTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 246

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| |||
Db        247 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 306

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||||  |||||||  |||||||||| | |||||   ||  |||| |||||||
Db        307 GAAGATTTTGCAACTTACTACTGTCAACAGTATTACAGTACTCCATTCACTTTCGGCCCT 366

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||   | ||||  ||  ||
Db        367 GGGACCAAAGTGGAAATCAAA 387


RESULT 5
US-09-240-274-216
; Sequence 216, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
;  APPLICANT: Siegel, Donald L.
;  TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
;  TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
;  FILE REFERENCE: 09596-42U2
;  CURRENT APPLICATION NUMBER: US/09/240,274
;  CURRENT FILING DATE: 1999-01-29
;  EARLIER APPLICATION NUMBER: 60/081,380
;  EARLIER FILING DATE: 1998-04-10
;  EARLIER APPLICATION NUMBER: 60/028,550
;  EARLIER FILING DATE: 1996-10-11
;  NUMBER OF SEQ ID NOS: 224
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 216
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: anti-Rh(D) antibody clone SH47
US-09-240-274-216

  Query Match            69.8%;  Score 224;  DB 3;  Length 321;
  Best Local Similarity  82.4%;  Pred. No. 7.7e-69;
  Matches  257;  Conservative    0;  Mismatches   55;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
              |||||||||||||||||||||||||| |||||||||||||||||||||||||||||| |
Db         63 TTGCCGGGCAAGTCAGAGCATTAGCAACTATTTAAATTGGTATCAGCAGAAACCAGGAAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
               | ||||| || || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAACCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| ||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| |||||||  |||||||||  | ||||||  ||   ||| ||||||||||||||
Db        243 TTTTGCAACTTACTACTGTCAACAGAGTTACAGTTATCCTCGCACGTTCGGCCAAGGGAC 302

Qy        306 ACGACTGGACAT 317
              |||| ||
Db        303 CAAGGTGGAGAT 314


RESULT 6
US-09-240-274-102
; Sequence 102, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
;  APPLICANT: Siegel, Donald L.
;  TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
;  TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
;  FILE REFERENCE: 09596-42U2
;  CURRENT APPLICATION NUMBER: US/09/240,274
;  CURRENT FILING DATE: 1999-01-29
;  EARLIER APPLICATION NUMBER: 60/081,380
;  EARLIER FILING DATE: 1998-04-10
;  EARLIER APPLICATION NUMBER: 60/028,550
;  EARLIER FILING DATE: 1996-10-11
;  NUMBER OF SEQ ID NOS: 224
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 102
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: anti-Rh(D) chain I02
US-09-240-274-102

  Query Match            69.5%;  Score 223.2;  DB 3;  Length 321;
  Best Local Similarity  81.6%;  Pred. No. 1.5e-68;
  Matches  258;  Conservative    0;  Mismatches   58;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 125
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||  |
Db         63 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
               | |||||||| || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| ||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| |||||||  |||||||||  | |||||   | |   || ||||||||||||||
Db        243 TTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCTGTGGACGTTCGGCCAAGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
              |||| ||  ||
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Db        303 CAAGGTGGAAATCAAA 318


RESULT 7
US-09-240-274-109
; Sequence 109, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
;  APPLICANT: Siegel, Donald L.
;  TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
;  TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
;  FILE REFERENCE: 09596-42U2
;  CURRENT APPLICATION NUMBER: US/09/240,274
;  CURRENT FILING DATE: 1999-01-29
;  EARLIER APPLICATION NUMBER: 60/081,380
;  EARLIER FILING DATE: 1998-04-10
;  EARLIER APPLICATION NUMBER: 60/028,550
;  EARLIER FILING DATE: 1996-10-11
;  NUMBER OF SEQ ID NOS: 224
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 109
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: anti-Rh(D) chain I09
US-09-240-274-109

  Query Match             69.5%;  Score 223.2;  DB 3;  Length 321;
  Best Local Similarity   81.6%;  Pred. No. 1.5e-68;
  Matches  258;  Conservative    0;  Mismatches   58;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  |
Db         63 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
                | ||||||| || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| ||||||||||||||||| ||||| ||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTCCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| ||||| |  ||||||||  | | ||||  |||  ||| || ||||| |||||
Db        243 TTTTGCAACTTATTACTGTCAACAGCTTAATAGTTACCCGTACACTTTTGGCCAGGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
                  ||||| ||  ||
Db        303 CAAGCTGGAGATCAAA 318


RESULT 8
US-09-240-274-215
; Sequence 215, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
;  APPLICANT: Siegel, Donald L.
;  TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
;  TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
;  FILE REFERENCE: 09596-42U2
;  CURRENT APPLICATION NUMBER: US/09/240,274
;  CURRENT FILING DATE: 1999-01-29
;  EARLIER APPLICATION NUMBER: 60/081,380
;  EARLIER FILING DATE: 1998-04-10
;  EARLIER APPLICATION NUMBER: 60/028,550
;  EARLIER FILING DATE: 1996-10-11
;  NUMBER OF SEQ ID NOS: 224
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 215
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: anti-Rh(D) antibody clone SH46
US-09-240-274-215

  Query Match             69.5%;  Score 223.2;  DB 3;  Length 321;
  Best Local Similarity   81.6%;  Pred. No. 1.5e-68;
  Matches  258;  Conservative    0;  Mismatches   58;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
              ||||||||||||||||  ||||||||||||||||||||||||||||||||||||||  |
Db         63 TTGCCGGGCAAGTCAGTACATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
                | ||||| || || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAATCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| ||||||||||||||||| ||||| ||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| ||||||| |||||||||| || ||||||  || | ||| |||||||  |||||
Db        243 TTTTGCAACTTACTACTGTCAACAGACTTACAGTTCCCCTAGCACTTTCGGCCCTGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
                  | |||| ||  ||
Db        303 CAAAGTGGATATCAAA 318


RESULT 9
US-09-240-274-217
; Sequence 217, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
;  APPLICANT: Siegel, Donald L.
;  TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
;  TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
;  FILE REFERENCE: 09596-42U2
;  CURRENT APPLICATION NUMBER: US/09/240,274
;  CURRENT FILING DATE: 1999-01-29
;  EARLIER APPLICATION NUMBER: 60/081,380
;  EARLIER FILING DATE: 1998-04-10
;  EARLIER APPLICATION NUMBER: 60/028,550
;  EARLIER FILING DATE: 1996-10-11
;  NUMBER OF SEQ ID NOS: 224
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 217
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: anti-Rh(D) antibody clone SH48
US-09-240-274-217

  Query Match             69.5%;  Score 223.2;  DB 3;  Length 321;
  Best Local Similarity   81.6%;  Pred. No. 1.5e-68;
  Matches  258;  Conservative    0;  Mismatches   58;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
              ||||||||||||||||  ||||||||||||||||||||||||||||||||||||||  |
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Db         63 TTGCCGGGCAAGTCAGTACATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
              | ||||| || || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAATCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| ||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| |||||||  ||||||||| || ||||||  || | ||| |||||||  |||||
Db        243 TTTTGCAACTTACTACTGTCAACAGACTTACAGTTCCCCTAGCACTTTCGGCCCTGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
                | |||| ||  ||
Db        303 CAAAGTGGATATCAAA 318


RESULT 10
US-09-240-274-218
; Sequence 218, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
;  APPLICANT: Siegel, Donald L.
;  TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
;  TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
;  FILE REFERENCE: 09596-42U2
;  CURRENT APPLICATION NUMBER: US/09/240,274
;  CURRENT FILING DATE: 1999-01-29
;  EARLIER APPLICATION NUMBER: 60/081,380
;  EARLIER FILING DATE: 1998-04-10
;  EARLIER APPLICATION NUMBER: 60/028,550
;  EARLIER FILING DATE: 1996-10-11
;  NUMBER OF SEQ ID NOS: 224
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 218
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: anti-Rh(D) antibody clone SH49
US-09-240-274-218

  Query Match             69.5%;  Score 223.2;  DB 3;  Length 321;
  Best Local Similarity   81.6%;  Pred. No. 1.5e-68;
  Matches  258;  Conservative    0;  Mismatches   58;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||| ||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCGTCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||   |
Db         63 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
              | |||||||| || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| ||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| |||||||  |||||||||  | |||||    |||    || ||||||||||||||
Db        243 TTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGTGGACGTTCGGCCAAGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
                  |||| ||  ||
Db        303 CAAGGTGGAAATCAAA 318


RESULT 11
US-09-859-053-29
; Sequence 29, Application US/09859053
; Patent No. 6803039
; GENERAL INFORMATION:
;  APPLICANT: Tsuji, Takashi
;  APPLICANT:  Tezuka, Katsunari
;  APPLICANT:  Hori, No. 6803039uaki
;  TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODY AGAINST A
;  TITLE OF INVENTION:  COSTIMULATORY SIGNAL TRANSDUCTION MOLECULE AILIM AND
;  TITLE OF INVENTION:  PHARMACEUTICAL USE THEREOF
;  FILE REFERENCE: 06501-079001
;  CURRENT APPLICATION NUMBER: US/09/859,053
;  CURRENT FILING DATE:  2001-05-16
;  PRIOR APPLICATION NUMBER: JP 2001-99508
;  PRIOR FILING DATE: 2001-03-30
;  PRIOR APPLICATION NUMBER: JP 2000-147116
;  PRIOR FILING DATE: 2000-05-18
;  NUMBER OF SEQ ID NOS: 43
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 29
;   LENGTH: 974
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: 5'UTR
;   LOCATION: (1)...(38)
;   NAME/KEY: CDS
;   LOCATION: (39)...(746)
;   NAME/KEY: 3'UTR
;   LOCATION: (750)...(974)
;   NAME/KEY: sig_peptide
;   LOCATION: (39)...(104)
US-09-859-053-29

  Query Match             69.1%;  Score 221.8;  DB 4;  Length 974;
  Best Local Similarity   80.7%;  Pred. No. 8.3e-68;
  Matches  259;  Conservative    0;  Mismatches   62;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||| ||| |||||||||||||||||||||||||||
Db        105 GACATCCAGATGACCCAGTCTCCATCTTCCGTGTCTGCATCTGTAGGAGACAGAGTCACC 164

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||||| |||||| | |||||||| |  |||   ||||||||||||||||||
Db        165 ATCACTTGTCGGGCGAGTCAGGGTATTAGCAGGTTGTTAGCCTGGTATCAGCAGAAACCA 224

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||| || || || ||    ||||| |    | ||   ||||||||
Db        225 GGGAAAGCCCCTAAACTCCTGATCTATGTTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 284

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | |||||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||
Db        285 AGGTTCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCT 344

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| ||||||| |||||||||| || ||||||| |||   || |||||||||
Db        345 GAAGATTTTGCAACTTACTATTGTCAACAGGCTAACAGTTTCCCGTGGACGTTCGGCCAA 404

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||     |||| ||  ||
Db        405 GGGACCAAGGTGGAAATCAAA 425


RESULT 12
US-09-240-274-199
; Sequence 199, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
;  APPLICANT: Siegel, Donald L.
;  TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
;  TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
;  FILE REFERENCE: 09596-42U2
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;  CURRENT APPLICATION NUMBER: US/09/240,274
;  CURRENT FILING DATE: 1999-01-29
;  EARLIER APPLICATION NUMBER: 60/081,380
;  EARLIER FILING DATE: 1998-04-10
;  EARLIER APPLICATION NUMBER: 60/028,550
;  EARLIER FILING DATE: 1996-10-11
;  NUMBER OF SEQ ID NOS: 224
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 199
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: anti-Rh(D) antibody clone SH13
US-09-240-274-199

  Query Match             69.0%;  Score 221.6;  DB 3;  Length 321;
  Best Local Similarity   81.3%;  Pred. No. 5.4e-68;
  Matches  257;  Conservative    0;  Mismatches   59;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  |
Db         63 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
                | |||||||| || || ||    ||||| |    |  |   ||||||||     | ||
Db        123 AGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCGAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| ||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              ||||||| |||||||  |||||||||  | |||||   ||   ||| || ||||| |||||
Db        243 TTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCCTACACTTTTGGCCAGGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
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Db        303 CAAGCTGGAGATCAAA 318


RESULT 13
US-09-042-353-358
; Sequence 358, Application US/09042353
; Patent No. 6255458
;  GENERAL INFORMATION:
;    APPLICANT:  Lonberg, Nils
;    APPLICANT:  Kay, Robert M.
;    TITLE OF INVENTION:  Transgenic No. 6255458-Human Animals for
;    TITLE OF INVENTION:  Producing Heterologous Antibodies
;    NUMBER OF SEQUENCES:  421
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Townsend and Townsend and Crew LLP
;      STREET:  Two Embarcadero Center, Eighth Floor
;      CITY:  San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94111-3834
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/042,353
;      FILING DATE:  13-MAR-1998
;      CLASSIFICATION:  800
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/810,279
;      FILING DATE:  17-DEC-1991
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/853,408
;      FILING DATE:  18-MAR-1992
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/904,068
;      FILING DATE:  23-JUN-1992
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/990,860
;      FILING DATE:  16-DEC-1992
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/053,131
;      FILING DATE:  26-APR-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/096,762
;      FILING DATE:  22-JUL-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/155,301
;      FILING DATE:  18-NOV-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/161,739
;      FILING DATE:  03-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/165,699
;      FILING DATE:  10-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/209,741
;      FILING DATE:  09-MAR-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/352,322
;      FILING DATE:  07-DEC-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/544,404
;      FILING DATE:  10-OCT-1995
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/728,463
;      FILING DATE:  10-OCT-1996
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  WO PCT/US96/16433
;      FILING DATE:  10-OCT-1996
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/758,417
;      FILING DATE:  02-DEC-1996
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  WO PCT/US97/21803
;      FILING DATE:  01-DEC-1997
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Apple, Randolph T.
;      REGISTRATION NUMBER:  36,429
;      REFERENCE/DOCKET NUMBER:  014643-009040US
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (415) 576-0200
;      TELEFAX:  (415) 576-0300
;  INFORMATION FOR SEQ ID NO:  358:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  388 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA
US-09-042-353-358

  Query Match             68.6%;  Score 220.2;  DB 3;  Length 388;
  Best Local Similarity   80.4%;  Pred. No. 1.9e-67;
  Matches  258;  Conservative    0;  Mismatches   63;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
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Db         67 GACATCCAGATGACCCAGTCTCCATCTTCCGTGTCTGCATCTGTAGGAGACAGAGTCACC 126

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||||| ||||||   |||||||||| |||   |||||||||| ||||||
Db        127 ATCACTTGTCGGGCGAGTCAGGATATTAGCAGCTGGTTAGCCTGGTATCAGCATAAACCA 186
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Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
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Db        187 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 246

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Db        247 AGGTTCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCT 306

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| ||||||| |||||||||| || | ||||| |||  ||| || |||||
Db        307 GAAGATTTTGCAACTTACTATTGTCAACAGGCTAATAGTTTCCCGTACACTTTTGGCCAG 366

Qy        301 GGGACACGACTGGACATTCAA 321
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Db        367 GGGACCAAGCTGGAGATCAAA 387


RESULT 14
US-08-758-417A-206
; Sequence 206, Application US/08758417A
; Patent No. 6300129
;  GENERAL INFORMATION:
;       APPLICANT: Lonberg, Nils
;                  Kay, Robert M.
;       TITLE OF INVENTION: Transgenic No. 6300129-Human Animals for
;                           Producing Heterologous Antibodies
;       NUMBER OF SEQUENCES: 417
;       CORRESPONDENCE ADDRESS:
;            ADDRESSEE: Townsend and Townsend and Crew LLP
;            STREET: Two Embarcadero Center, Eighth Floor
;            CITY: San Francisco
;            STATE: California
;            COUNTRY: USA
;            ZIP: 94111-3834
;       COMPUTER READABLE FORM:
;            MEDIUM TYPE: Floppy disk
;            COMPUTER: IBM PC compatible
;            OPERATING SYSTEM: PC-DOS/MS-DOS
;            SOFTWARE: PatentIn Release #1.0, Version #1.30
;       CURRENT APPLICATION DATA:
;            APPLICATION NUMBER: US/08/758,417A
;            FILING DATE: 02-Dec-1996
;            CLASSIFICATION: <Unknown>
;       PRIOR APPLICATION DATA:
;            APPLICATION NUMBER: US 08/728,463
;            FILING DATE: 10-OCT-1996
;            APPLICATION NUMBER: US 08/544,404
;            FILING DATE: 10-OCT-1995
;            APPLICATION NUMBER: US 08/352,322
;            FILING DATE: 07-DEC-1994
;            APPLICATION NUMBER: US 08/209,741
;            FILING DATE: 09-MAR-1994
;            APPLICATION NUMBER: US 08/165,699
;            FILING DATE: 10-DEC-1993
;            APPLICATION NUMBER: US 08/161,739
;            FILING DATE: 03-DEC-1993
;            APPLICATION NUMBER: US 08/155,301
;            FILING DATE: 18-NOV-1993
;            APPLICATION NUMBER: US 08/096,762
;            FILING DATE: 22-JUL-1993
;            APPLICATION NUMBER: US 08/053,131
;            FILING DATE: 26-APR-1993
;            APPLICATION NUMBER: US 07/990,860
;            FILING DATE: 16-DEC-1992
;       ATTORNEY/AGENT INFORMATION:
;            NAME: Serafini, Andrew T.
;            REGISTRATION NUMBER: 41,303
;            REFERENCE/DOCKET NUMBER: 014643-009030US
;       TELECOMMUNICATION INFORMATION:
;            TELEPHONE: (415) 576-0200
;            TELEFAX: (415) 576-0300
;  INFORMATION FOR SEQ ID NO: 206:
;       SEQUENCE CHARACTERISTICS:
;            LENGTH: 388 base pairs
;            TYPE: nucleic acid
;            STRANDEDNESS: single
;            TOPOLOGY: linear
;       MOLECULE TYPE: DNA
;       SEQUENCE DESCRIPTION: SEQ ID NO: 206:
US-08-758-417A-206

  Query Match           68.6%;  Score 220.2;  DB 3;  Length 388;
  Best Local Similarity   80.4%;  Pred. No. 1.9e-67;
  Matches  258;  Conservative    0;  Mismatches   63;  Indels    0;  Gaps    0;

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Db         67 GACATCCAGATGACCCAGTCTCCATCTTCCGTGTCTGCATCTGTAGGAGACAGAGTCACC 126

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
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Db        127 ATCACTTGTCGGGCGAGTCAGGATATTAGCAGCTGGTTAGCCTGGTATCAGCATAAACCA 186

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
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Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
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Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
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Qy        301 GGGACACGACTGGACATTCAA 321
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Db        367 GGGACCAAGCTGGAGATCAAA 387


RESULT 15
US-09-240-274-107
; Sequence 107, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
;  APPLICANT: Siegel, Donald L.
;  TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
;  TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
;  FILE REFERENCE: 09596-42U2
;  CURRENT APPLICATION NUMBER: US/09/240,274
;  CURRENT FILING DATE: 1999-01-29
;  EARLIER APPLICATION NUMBER: 60/081,380
;  EARLIER FILING DATE: 1998-04-10
;  EARLIER APPLICATION NUMBER: 60/028,550
;  EARLIER FILING DATE: 1996-10-11
;  NUMBER OF SEQ ID NOS: 224
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 107
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: anti-Rh(D) chain I07
US-09-240-274-107

  Query Match           68.5%;  Score 220;  DB 3;  Length 321;
  Best Local Similarity   81.0%;  Pred. No. 2e-67;
  Matches  256;  Conservative    0;  Mismatches   60;  Indels    0;  Gaps    0;

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              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
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              | ||||||||| || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| ||||||||||||||||| ||||| ||||||||
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Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              |||||| ||||||| ||||||||| | |||||   ||    || ||||||  ||||||
Db        243 TTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCTCGAACTTTCGGCGGAGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
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Db        303 CAAGGTGGAGATCAAA 318
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GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

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Title: US-09-403-107-147
Perfect score: 321
Sequence: 1 gagctccagatgacccagtc.....ggacacgactggacattcaa 321

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

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Maximum Match 100%
Listing first 45 summaries

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26: /cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	321	100.0	321	15	US-10-325-694-147	Sequence 147, App
2	255.4	79.6	321	15	US-10-325-694-141	Sequence 141, App
3	245.8	76.6	324	19	US-10-344-514-7	Sequence 7, Appli
4	245.8	76.6	324	19	US-10-344-514-8	Sequence 8, Appli
5	240.8	75.0	321	10	US-09-848-798-106	Sequence 106, App
6	240.2	74.8	321	19	US-10-703-714-11	Sequence 11, Appl
7	240.2	74.8	321	19	US-10-703-714-19	Sequence 19, Appl
8	239.4	74.6	322	17	US-10-309-762-215	Sequence 215, App
9	239.4	74.6	322	17	US-10-309-762-228	Sequence 228, App
10	237.8	74.1	321	21	US-10-727-155-155	Sequence 155, App
11	236.2	73.6	321	21	US-10-727-155-141	Sequence 141, App
12	236.2	73.6	321	21	US-10-727-155-159	Sequence 159, App
13	236.2	73.6	322	17	US-10-309-762-213	Sequence 213, App
14	236.2	73.6	322	17	US-10-309-762-238	Sequence 238, App
15	236.2	73.6	458	21	US-10-644-277-55	Sequence 55, Appl
16	235	73.2	1106	17	US-10-264-049-121	Sequence 121, App
17	233	72.6	321	21	US-10-727-155-149	Sequence 149, App
18	233	72.6	321	21	US-10-949-135-25	Sequence 25, Appl
19	231.4	72.1	333	16	US-10-203-754A-60	Sequence 60, Appl
20	231.4	72.1	720	9	US-09-192-854-1	Sequence 1, Appli
21	231.4	72.1	720	9	US-09-968-561A-1	Sequence 1, Appli
22	231.4	72.1	720	10	US-09-968-744A-1	Sequence 1, Appli
23	231.4	72.1	720	11	US-09-968-561A-1	Sequence 1, Appli
24	231.4	72.1	720	20	US-10-744-774-2	Sequence 2, Appli
25	231.4	72.1	900	16	US-10-203-754A-64	Sequence 64, Appl
26	230.4	71.8	321	21	US-10-949-135-9	Sequence 9, Appli
27	229.8	71.6	321	17	US-10-338-366-11	Sequence 11, Appl
28	229.8	71.6	322	22	US-10-916-840-115	Sequence 115, App
29	229.8	71.6	324	19	US-10-344-514-3	Sequence 3, Appli
30	229.8	71.6	324	19	US-10-344-514-4	Sequence 4, Appli
31	229.8	71.6	729	15	US-10-216-484-125	Sequence 125, App
32	229.8	71.6	729	16	US-10-384-933-125	Sequence 125, App
33	229.6	71.5	348	22	US-10-916-758-40	Sequence 40, Appl
34	228.2	71.1	322	17	US-10-309-762-226	Sequence 226, App
35	228.2	71.1	322	17	US-10-309-762-227	Sequence 227, App
36	228.2	71.1	324	20	US-10-409-814A-3	Sequence 3, Appli
37	228.2	71.1	405	21	US-10-783-311-15	Sequence 15, Appl
38	228.2	71.1	405	21	US-10-783-311-16	Sequence 16, Appl
39	228.2	71.1	702	21	US-10-938-353-3	Sequence 3, Appli
40	228.2	71.1	714	14	US-10-153-382-18	Sequence 18, Appl
41	228.2	71.1	714	20	US-10-612-497-62	Sequence 62, Appl
42	228.2	71.1	714	20	US-10-776-649-62	Sequence 62, Appl
43	228.2	71.1	819	14	US-10-158-646-65	Sequence 65, Appl
44	226.6	70.6	322	16	US-10-041-860-74	Sequence 74, Appl
45	226.6	70.6	322	17	US-10-309-762-221	Sequence 221, App

ALIGNMENTS

RESULT 1
US-10-325-694-147
; Sequence 147, Application US/10325694
; Publication No. US20030148463A1
; GENERAL INFORMATION:
; APPLICANT: KUFER, PETER
; APPLICANT: RAUM, TOBIAS
; TITLE OF INVENTION: NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN
; TITLE OF INVENTION: RECEPTORS AND USES THEREOF
; FILE REFERENCE: 38164000
; CURRENT APPLICATION NUMBER: US/10/325,694
; CURRENT FILING DATE: 2002-12-19
; PRIOR APPLICATION NUMBER: US/09/403,107
; PRIOR FILING DATE: 1999-10-14
; NUMBER OF SEQ ID NOS: 152
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 147
; LENGTH: 321
; TYPE: DNA
; ORGANISM: HUMAN
US-10-325-694-147

Query Match 100.0%; Score 321; DB 15; Length 321;
Best Local Similarity 100.0%; Pred. No. 3.3e-101;
Matches 321; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT 2
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; Sequence 141, Application US/10325694
; Publication No. US20030148463A1
; GENERAL INFORMATION:
;  APPLICANT: KUFER, PETER
;  APPLICANT:  RAUM, TOBIAS
;  TITLE OF INVENTION: NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN
;  TITLE OF INVENTION:  RECEPTORS AND USES THEREOF
;  FILE REFERENCE: 38164000
;  CURRENT APPLICATION NUMBER: US/10/325,694
;  CURRENT FILING DATE:  2002-12-19
;  PRIOR APPLICATION NUMBER: US/09/403,107
;  PRIOR FILING DATE: 1999-10-14
;  NUMBER OF SEQ ID NOS: 152
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 141
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: HUMAN
US-10-325-694-141

  Query Match             79.6%;  Score 255.4;  DB 15;  Length 321;
  Best Local Similarity   87.2%;  Pred. No. 2.5e-78;
  Matches  280;  Conservative    0;  Mismatches   41;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||||||||||||||||||||||||||||||||||| ||||| ||||||||||||||||
Db          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCTTCTGTGGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||| ||| |||||||||||||||||||||||||| ||||||||||||||||||||
Db         61 ATCACTTGTCGGACAAGTCAGAGCATTAGCAGCTATTAAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||||||||||||||||||||||||||||||  ||||||||||||||||||||||||||||
Db        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGCGATCTACCCGGGAATCCGGGGTCCCTGAC 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              |||||||| ||||| |  ||||||||| ||| |||||||||||||||||| || || |||
Db        181 CGATTCAGTGGCAGCGGGTCTGGGACAGATTTCACTCTCACCATCAGCAGTCTACAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||| ||| |||||||  ||||| |||  | ||    | |||  ||| || |||||
Db        241 GAAGATTCTGCAACTTACTACTGTCAGCAGAGTTACGACATCCCGTACACTTTTGGCCAG 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||    ||||| ||  ||
Db        301 GGGACCAAGCTGGAGATCAAA 321


RESULT 3
US-10-344-514-7
; Sequence 7, Application US/10344514
; Publication No. US20040120951A1
; GENERAL INFORMATION:
;  APPLICANT: NAKASHIMA, Toshihiro et al.
;  TITLE OF INVENTION: HUMAN ANTI-FACTOR VIII ANTIBODY
;  FILE REFERENCE: 0020-5111P
;  CURRENT APPLICATION NUMBER: US/10/344,514
;  CURRENT FILING DATE:  2003-07-28
;  PRIOR APPLICATION NUMBER: JP 2001-177640
;  PRIOR FILING DATE: 2001-06-12
;  NUMBER OF SEQ ID NOS: 18
; SEQ ID NO 7
;   LENGTH: 324
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-344-514-7

  Query Match             76.6%;  Score 245.8;  DB 19;  Length 324;
  Best Local Similarity   85.4%;  Pred. No. 5.6e-75;
  Matches  274;  Conservative    0;  Mismatches   47;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||||| || || ||    ||||| |    | ||   ||||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| ||||||| |||||||||| ||| |||||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||||  | |||||    ||||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 4
US-10-344-514-8
; Sequence 8, Application US/10344514
; Publication No. US20040120951A1
; GENERAL INFORMATION:
;  APPLICANT: NAKASHIMA, Toshihiro et al.
;  TITLE OF INVENTION: HUMAN ANTI-FACTOR VIII ANTIBODY
;  FILE REFERENCE: 0020-5111P
;  CURRENT APPLICATION NUMBER: US/10/344,514
;  CURRENT FILING DATE:  2003-07-28
;  PRIOR APPLICATION NUMBER: JP 2001-177640
;  PRIOR FILING DATE: 2001-06-12
;  NUMBER OF SEQ ID NOS: 18
; SEQ ID NO 8
;   LENGTH: 324
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)..(324)
US-10-344-514-8

  Query Match             76.6%;  Score 245.8;  DB 19;  Length 324;
  Best Local Similarity   85.4%;  Pred. No. 5.6e-75;
  Matches  274;  Conservative    0;  Mismatches   47;  Indels    0;  Gaps    0;
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Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||||| || || ||    ||||| |    | ||   ||||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||| ||||||| |||||||||| ||| ||||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||| | |||||   ||||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 5
US-09-848-798-106
; Sequence 106, Application US/09848798
; Publication No. US20030040605A1
; GENERAL INFORMATION:
;  APPLICANT: Siegel, Donald L.
;  TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
;  TITLE OF INVENTION:  SORTING METHOD FOR PRODUCTION THEREOF
;  FILE REFERENCE: 09596-42U2
;  CURRENT APPLICATION NUMBER: US/09/848,798
;  CURRENT FILING DATE:  2001-05-04
;  PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,274
;  PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-29
;  PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550
;  PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11
;  NUMBER OF SEQ ID NOS: 224
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 106
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: anti-Rh(D) chain I06
US-09-848-798-106

  Query Match             75.0%;  Score 240.8;  DB 10;  Length 321;
  Best Local Similarity   85.1%;  Pred. No. 3.1e-73;
  Matches  269;  Conservative    0;  Mismatches   47;  Indels    0;  Gaps    0;

Qy          6 CCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 65
              | || | ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          3 CGAGCTCACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAC 62

Qy         66 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGACA 125
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  |
Db         63 TTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGGAA 122

Qy        126 GCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGATT 185
              | |||||||| || || ||    ||||| |    | ||   ||||||||     | ||
Db        123 AGCCCCTAAGCTCCTGATCTATGCCGCATCCAGTTTGCAAAGTGGGGTCCCATCAAGGTT 182

Qy        186 CAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAGA 245
              ||| ||||||| |||||||||| ||| ||||||||||||||||| ||||| |||||||||
Db        183 CAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCTGAAGA 242

Qy        246 TTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGAC 305
              ||||||  ||||||| |||||||| | |||||   ||||||||||||||||||||||
Db        243 TTTTGCAACTTACTACTGTCAACAGAGTTACAGTACCCCGATCACCTTCGGCCAAGGGAC 302

Qy        306 ACGACTGGACATTCAA 321
              |||||||||  ||| ||
Db        303 ACGACTGGAGATTAAA 318


RESULT 6
US-10-703-714-11
; Sequence 11, Application US/10703714
; Publication No. US20040170630A1
; GENERAL INFORMATION:
;  APPLICANT: Huang, Haichun
;  APPLICANT:  Holmes, Steven
;  APPLICANT:  Mason, Sean
;  TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO HEPARANASE
;  FILE REFERENCE: MXI-294
;  CURRENT APPLICATION NUMBER: US/10/703,714
;  CURRENT FILING DATE:  2003-11-07
;  PRIOR APPLICATION NUMBER: 60/424803
;  PRIOR FILING DATE: 2002-11-07
;  NUMBER OF SEQ ID NOS: 64
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 11
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1)...(321)
US-10-703-714-11

  Query Match             74.8%;  Score 240.2;  DB 19;  Length 321;
  Best Local Similarity   84.9%;  Pred. No. 5e-73;
  Matches  269;  Conservative    0;  Mismatches   48;  Indels    0;  Gaps    0;

Qy          5 TCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64
              ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          5 TCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64

Qy         65 CTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGAC 124
              |||||||||||||||| ||||||||||   ||||   ||||||||||||||||||||
Db         65 CTTGCCGGGCAAGTCAGGGCATTAGCAGTGCTTTAGCCTGGTATCAGCAGAAACCAGGGA 124

Qy        125 AGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGAT 184
              |  |||||||||| || || ||    || || |    ||||   ||||||||     | |
Db        125 AAGCTCCTAAGCTCCTGATCTATGATGCCTCCAGTTTGGAAAGTGGGGTCCCATCAAGGT 184

Qy        185 TCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244
              |||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||||||
Db        185 TCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244

Qy        245 ATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGA 304
              ||||||| ||||| |  |||||||||| | | ||||  ||||||||||||||||||||
Db        245 ATTTTGCAACTTATTACTGTCAACAGTTTAATAGTTACCCGATCACCTTCGGCCAAGGGA 304

Qy        305 CACGACTGGACATTCAA 321
              ||||||||||  ||| ||
Db        305 CACGACTGGAGATTAAA 321


RESULT 7
US-10-703-714-19
; Sequence 19, Application US/10703714
; Publication No. US20040170630A1
; GENERAL INFORMATION:
;  APPLICANT: Huang, Haichun
;  APPLICANT:  Holmes, Steven
;  APPLICANT:  Mason, Sean
;  TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO HEPARANASE
;  FILE REFERENCE: MXI-294
;  CURRENT APPLICATION NUMBER: US/10/703,714
;  CURRENT FILING DATE:  2003-11-07
;  PRIOR APPLICATION NUMBER: 60/424803
```

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;  PRIOR FILING DATE: 2002-11-07
;  NUMBER OF SEQ ID NOS: 64
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 19
;    LENGTH: 321
;    TYPE: DNA
;    ORGANISM: Homo sapiens
;    FEATURE:
;    NAME/KEY: CDS
;    LOCATION: (1)...(321)
US-10-703-714-19

  Query Match            74.8%;  Score 240.2;  DB 19;  Length 321;
  Best Local Similarity  84.9%;  Pred. No. 5e-73;
  Matches  269;  Conservative    0;  Mismatches   48;  Indels    0;  Gaps    0;

Qy          5 TCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64
              ||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          5 TCCAGTTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCA 64

Qy         65 CTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCAGGAC 124
              |||||||||||||||| ||||||||||   ||||   ||||||||||||||||||||
Db         65 CTTGCCGGGCAAGTCAGGGCATTAGCAGTGCTTTAGCCTGGTATCAGCAGAAACCAGGGA 124

Qy        125 AGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGACCGAT 184
              |  |||||||||| || || ||    || || |    ||||   ||||||||     | |
Db        125 AAGCTCCTAAGCTCCTGATCTATGATGCCTCCAGTTTGGAAAGTGGGGTCCCATCAAGGT 184

Qy        185 TCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244
              |||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||||
Db        185 TCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAG 244

Qy        245 ATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAAGGGA 304
              ||||||| ||||| |  |||||||||| | | ||||  ||||||||||||||||||||
Db        245 ATTTTGCAACTTATTACTGTCAACAGTTTAATAGTTACCCGATCACCTTCGGCCAAGGGA 304

Qy        305 CACGACTGGACATTCAA 321
              |||||||||| ||| ||
Db        305 CACGACTGGAGATTAAA 321


RESULT 8
US-10-309-762-215
; Sequence 215, Application US/10309762
; Publication No. US20040018198A1
; GENERAL INFORMATION:
;  APPLICANT: Gudas, Jean
;  APPLICANT:  Foltz, Ian
;  APPLICANT:  Handa, Masahisa
;  APPLICANT:  Gallo, Michael
;  TITLE OF INVENTION: ANTIBODIES AGAINST CARBOXIC ANHYDRASE IX
;  TITLE OF INVENTION:  (CA IX) TUMOR ANTIGEN
;  FILE REFERENCE: ABGENIX.027A
;  CURRENT APPLICATION NUMBER: US/10/309,762
;  CURRENT FILING DATE:  2002-12-02
;  PRIOR APPLICATION NUMBER: 60/337275
;  PRIOR FILING DATE: 2001-12-03
;  NUMBER OF SEQ ID NOS: 246
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 215
;    LENGTH: 322
;    TYPE: DNA
;    ORGANISM: Homo sapiens
US-10-309-762-215

  Query Match            74.6%;  Score 239.4;  DB 17;  Length 322;
  Best Local Similarity  84.1%;  Pred. No. 9.5e-73;
  Matches  270;  Conservative    0;  Mismatches   51;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  |||||||||||||||||||| ||| ||||||||||||||||||||||||||||||
Db          1 GACATCCAGATGACCCAGTCTCCATCTTCCGTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||||| |||||| | ||||||| ||  ||||  ||||||||||||||||||
Db         61 ATCACTTGTCGGGCGAGTCAGGGTATTAGCATCTGGTTAACCTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||||| || || ||    |||||  |    | ||   ||||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||
Db        181 AGGTTCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| ||||||| |||||||||| || ||||||| || ||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTATTGTCAACAGGCTAACAGTTTCCCTATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 9
US-10-309-762-228
; Sequence 228, Application US/10309762
; Publication No. US20040018198A1
; GENERAL INFORMATION:
;  APPLICANT: Gudas, Jean
;  APPLICANT:  Foltz, Ian
;  APPLICANT:  Handa, Masahisa
;  APPLICANT:  Gallo, Michael
;  TITLE OF INVENTION: ANTIBODIES AGAINST CARBOXIC ANHYDRASE IX
;  TITLE OF INVENTION:  (CA IX) TUMOR ANTIGEN
;  FILE REFERENCE: ABGENIX.027A
;  CURRENT APPLICATION NUMBER: US/10/309,762
;  CURRENT FILING DATE:  2002-12-02
;  PRIOR APPLICATION NUMBER: 60/337275
;  PRIOR FILING DATE: 2001-12-03
;  NUMBER OF SEQ ID NOS: 246
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 228
;    LENGTH: 322
;    TYPE: DNA
;    ORGANISM: Homo sapiens
US-10-309-762-228

  Query Match            74.6%;  Score 239.4;  DB 17;  Length 322;
  Best Local Similarity  84.1%;  Pred. No. 9.5e-73;
  Matches  270;  Conservative    0;  Mismatches   51;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  |||||||||||||||||||| ||| ||||||||||||||||||||||||||||||
Db          1 GACATCCAGATGACCCAGTCTCCATCTTCCGTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||||| |||||| | ||||||| ||  ||||  ||||||||||||||||||
Db         61 ATCACTTGTCGGGCGAGTCAGGGTATTAGCATCTGGTTAACCTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | |||||||| || || ||    |||||  |    | ||   ||||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
              | ||||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||
Db        181 AGGTTCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| ||||||| |||||||||| || ||||||| || ||||||||||||||||
Db        241 GAAGATTTTGCAACTTACTATTGTCAACAGGCTAACAGTTTCCCTATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321
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```
RESULT 10
US-10-727-155-155
; Sequence 155, Application US/10727155
; Publication No. US20050049402A1
; GENERAL INFORMATION:
;  APPLICANT: John S. Babcook
;  APPLICANT:  Jaspal S. Kang
;  APPLICANT:  Orit Foord
;  APPLICANT:  Larry Green
;  APPLICANT:  Xiao Feng
;  APPLICANT:  Scott Klakamp
;  APPLICANT:  Mary Haak-Frendscho
;  APPLICANT:  Palaniswami Rathanaswami
;  APPLICANT:  Craig Pigott
;  APPLICANT:  Meina Liang
;  APPLICANT:  Rozanne Lee
;  APPLICANT:  Kathy Manchulencho
;  APPLICANT:  Raffaella Faggioni
;  APPLICANT:  Giorgio Senaldi
;  APPLICANT:  Qiaojuan Jane Su
;  TITLE OF INVENTION: ANTIBODIES DIRECTED TO TUMOR NECROSIS
;  TITLE OF INVENTION:  FACTOR AND USES THEREOF
;  FILE REFERENCE: ABGENIX.073A
;  CURRENT APPLICATION NUMBER: US/10/727,155
;  CURRENT FILING DATE:  2003-12-02
;  PRIOR APPLICATION NUMBER: 60/430729
;  PRIOR FILING DATE: 2002-12-02
;  NUMBER OF SEQ ID NOS: 320
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 155
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-727-155-155

  Query Match             74.1%;  Score 237.8;  DB 21;  Length 321;
  Best Local Similarity   83.8%;  Pred. No. 3.4e-72;
  Matches  269;  Conservative    0;  Mismatches   52;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              ||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGACAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||| || | || || ||    ||||| |    | ||   ||||||||
Db        121 GGGAAAGCCCCTGAGGTCCTGATCTATGCTGCATCCAATTTGCAACGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| ||||||| |||||||||| ||| |||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||||  |  ||||   |  ||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTCCAGTACCCTCATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 11
US-10-727-155-141
; Sequence 141, Application US/10727155
; Publication No. US20050049402A1
; GENERAL INFORMATION:
;  APPLICANT: John S. Babcook
;  APPLICANT:  Jaspal S. Kang
;  APPLICANT:  Orit Foord
;  APPLICANT:  Larry Green
;  APPLICANT:  Xiao Feng
;  APPLICANT:  Scott Klakamp
;  APPLICANT:  Mary Haak-Frendscho
;  APPLICANT:  Palaniswami Rathanaswami
;  APPLICANT:  Craig Pigott
;  APPLICANT:  Meina Liang
;  APPLICANT:  Rozanne Lee
;  APPLICANT:  Kathy Manchulencho
;  APPLICANT:  Raffaella Faggioni
;  APPLICANT:  Giorgio Senaldi
;  APPLICANT:  Qiaojuan Jane Su
;  TITLE OF INVENTION: ANTIBODIES DIRECTED TO TUMOR NECROSIS
;  TITLE OF INVENTION:  FACTOR AND USES THEREOF
;  FILE REFERENCE: ABGENIX.073A
;  CURRENT APPLICATION NUMBER: US/10/727,155
;  CURRENT FILING DATE:  2003-12-02
;  PRIOR APPLICATION NUMBER: 60/430729
;  PRIOR FILING DATE: 2002-12-02
;  NUMBER OF SEQ ID NOS: 320
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 141
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-727-155-141

  Query Match             73.6%;  Score 236.2;  DB 21;  Length 321;
  Best Local Similarity   83.5%;  Pred. No. 1.2e-71;
  Matches  268;  Conservative    0;  Mismatches   53;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCGCC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||||| |||||||||||||||||| ||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGACAAGTCAGAGCATTAGCAGTTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||| |||| || || ||    ||||| |    | ||   ||||||||
Db        121 GGGAAAGCCCCTGAGCTCCTGATCTATGCTGCATCCAATTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| ||||||| |||||||||| ||| |||||||||||||||| ||||| |||
Db        181 AGGTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCAACCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||| |||||||  |||||||||  |  ||||   |  ||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTCCAGTACCCTCATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 12
US-10-727-155-159
; Sequence 159, Application US/10727155
; Publication No. US20050049402A1
; GENERAL INFORMATION:
;  APPLICANT: John S. Babcook
;  APPLICANT:  Jaspal S. Kang
;  APPLICANT:  Orit Foord
;  APPLICANT:  Larry Green
;  APPLICANT:  Xiao Feng
;  APPLICANT:  Scott Klakamp
;  APPLICANT:  Mary Haak-Frendscho
;  APPLICANT:  Palaniswami Rathanaswami
;  APPLICANT:  Craig Pigott
;  APPLICANT:  Meina Liang
;  APPLICANT:  Rozanne Lee
;  APPLICANT:  Kathy Manchulencho
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;  APPLICANT:  Raffaella Faggioni
;  APPLICANT:  Giorgio Senaldi
;  APPLICANT:  Qiaojuan Jane Su
;  TITLE OF INVENTION: ANTIBODIES DIRECTED TO TUMOR NECROSIS
;  TITLE OF INVENTION:  FACTOR AND USES THEREOF
;  FILE REFERENCE: ABGENIX.073A
;  CURRENT APPLICATION NUMBER: US/10/727,155
;  CURRENT FILING DATE:  2003-12-02
;  PRIOR APPLICATION NUMBER: 60/430729
;  PRIOR FILING DATE: 2002-12-02
;  NUMBER OF SEQ ID NOS: 320
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 159
;   LENGTH: 321
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-727-155-159

  Query Match             73.6%;  Score 236.2;  DB 21;  Length 321;
  Best Local Similarity   83.5%;  Pred. No. 1.2e-71;
  Matches  268;  Conservative    0;  Mismatches   53;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 ATCACTTGCCGGACAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||| | || || || ||    |||| ||    | ||   ||||||||
Db        121 GGGAAAGCCCCTGAACTCCTGATCTATGCTGCATTTAATTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               |  |||| |||||||| |||||||||| ||| |||||||||||||||||| ||||| |||
Db        181 AGGATCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGTCTGCACCCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              ||||||||||| |||||||  |||||||||  |  ||||   |  ||||||||||||||
Db        241 GAAGATTTTGCAACTTACTACTGTCAACAGAGTTCCAGTACCCTCATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 13
US-10-309-762-213
; Sequence 213, Application US/10309762
; Publication No. US20040018198A1
; GENERAL INFORMATION:
;  APPLICANT: Gudas, Jean
;  APPLICANT:  Foltz, Ian
;  APPLICANT:  Handa, Masahisa
;  APPLICANT:  Gallo, Michael
;  TITLE OF INVENTION: ANTIBODIES AGAINST CARBOXIC ANHYDRASE IX
;  TITLE OF INVENTION:  (CA IX) TUMOR ANTIGEN
;  FILE REFERENCE: ABGENIX.027A
;  CURRENT APPLICATION NUMBER: US/10/309,762
;  CURRENT FILING DATE:  2002-12-02
;  PRIOR APPLICATION NUMBER: 60/337275
;  PRIOR FILING DATE: 2001-12-03
;  NUMBER OF SEQ ID NOS: 246
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 213
;   LENGTH: 322
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-309-762-213

  Query Match             73.6%;  Score 236.2;  DB 17;  Length 322;
  Best Local Similarity   83.5%;  Pred. No. 1.2e-71;
  Matches  268;  Conservative    0;  Mismatches   53;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGATGACCCAGTCTCCATCTTCCGTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||||| |||||| | |||||||||| |||   |||| ||||||||||||||
Db         61 ATCACTTGTCGGGCGAGTCAGGGTATTAGCAGCTGGTTAGCCTGGTTTCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | || ||||| || || ||    ||||| |    | ||   ||||||||
Db        121 GGGAAAGCCCCCAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | |||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||||
Db        181 AGGTTCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||||| ||||||| |||||||||| || ||||||| || ||||||||||||||
Db        241 GAAGATTTTGCAACTTACTATTGTCAACAGGCTAACAGTTTCCCTATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              |||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 14
US-10-309-762-238
; Sequence 238, Application US/10309762
; Publication No. US20040018198A1
; GENERAL INFORMATION:
;  APPLICANT: Gudas, Jean
;  APPLICANT:  Foltz, Ian
;  APPLICANT:  Handa, Masahisa
;  APPLICANT:  Gallo, Michael
;  TITLE OF INVENTION: ANTIBODIES AGAINST CARBOXIC ANHYDRASE IX
;  TITLE OF INVENTION:  (CA IX) TUMOR ANTIGEN
;  FILE REFERENCE: ABGENIX.027A
;  CURRENT APPLICATION NUMBER: US/10/309,762
;  CURRENT FILING DATE:  2002-12-02
;  PRIOR APPLICATION NUMBER: 60/337275
;  PRIOR FILING DATE: 2001-12-03
;  NUMBER OF SEQ ID NOS: 246
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 238
;   LENGTH: 322
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-309-762-238

  Query Match             73.6%;  Score 236.2;  DB 17;  Length 322;
  Best Local Similarity   83.5%;  Pred. No. 1.2e-71;
  Matches  268;  Conservative    0;  Mismatches   53;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGATGACCCAGTCTCCATCTTCCGTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||| ||||| |||||| | |||||||||| |||   |||| ||||||||||||||
Db         61 ATCACTTGTCGGGCGAGTCAGGGTATTAGCAGCTGGTTAGCCTGGTTTCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | || ||||| || || ||    ||||| |    | ||   ||||||||
Db        121 GGGAAAGCCCCCAAGCTCCTGATCTATGCTGCATCCAGTTTGCAAAGTGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | |||||||||||| |||||||||| ||| ||||||||||||||||||||||||||||||
Db        181 AGGTTCAGCGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||||||||| ||||||| |||||||||| || ||||||| || ||||||||||||||
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Db        241 GAAGATTTTGCAACTTACTATTGTCAACAGGCTAACAGTTTCCCTATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


RESULT 15
US-10-644-277-55
; Sequence 55, Application US/10644277
; Publication No. US20050058639A1
; GENERAL INFORMATION:
;  APPLICANT: Gudas, Jean M.
;  APPLICANT:  Haak-Frendscho, Mary
;  APPLICANT:  Foord, Orit
;  APPLICANT:  Liang, Meina L.
;  APPLICANT:  Ahluwalia, Kiran
;  APPLICANT:  Bhakta, Sunil
;  TITLE OF INVENTION: ANTIBODIES DIRECTED TO MONOCYTE
;  TITLE OF INVENTION:  CHEMO-ATTRACTANT PROTEIN-1 (MCP-1) AND USES THEREOF
;  FILE REFERENCE: ABGENIX.091A
;  CURRENT APPLICATION NUMBER: US/10/644,277
;  CURRENT FILING DATE:  2003-08-19
;  PRIOR APPLICATION NUMBER: 60/404,802
;  PRIOR FILING DATE: 2002-08-19
;  NUMBER OF SEQ ID NOS: 149
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 55
;   LENGTH: 458
;   TYPE: DNA
;   ORGANISM: Homosapien
US-10-644-277-55

  Query Match             73.6%;  Score 236.2;  DB 21;  Length 458;
  Best Local Similarity   83.5%;  Pred. No. 1.4e-71;
  Matches  268;  Conservative    0;  Mismatches   53;  Indels    0;  Gaps    0;

Qy          1 GAGCTCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60
              ||  ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACC 60

Qy         61 ATCACTTGCCGGGCAAGTCAGAGCATTAGCAGCTATTTAAATTGGTATCAGCAGAAACCA 120
              |||||||||| ||| ||||||  ||||| || ||||||||||||||||||||||||||||
Db         61 ATCACTTGCCAGGCGAGTCAGGACATTACCACCTATTTAAATTGGTATCAGCAGAAACCA 120

Qy        121 GGACAGCCTCCTAAGCTGCTCATTTACTGGGCATCTACCCGGGAATCCGGGGTCCCTGAC 180
              ||  |  | ||||||||  || || |||   ||||| |    |||| | ||||||||
Db        121 GGGAAAGCCCCTAAGCTCCTGATCTACGATGCATCCAATTTGGAAACAGGGGTCCCATCA 180

Qy        181 CGATTCAGCGGCAGTGAATCTGGGACAAATTACACTCTCACCATCAGCAGCCTGCAGCCT 240
               | ||||| || |||| |||||||||| |||  ||| ||||||||||||||||||||||
Db        181 AGGTTCAGTGGAAGTGGATCTGGGACAGATTTTACTTTCACCATCAGCAGCCTGCAGCCT 240

Qy        241 GAAGATTTTGCTACTTACTTTTGTCAACAGTCTGACAGTTTGCCGATCACCTTCGGCCAA 300
              |||||| |||| || || |  |||||||| | ||| | | | ||||||||||||||||||
Db        241 GAAGATATTGCAACATATTACTGTCAACAATATGATAATCTCCCGATCACCTTCGGCCAA 300

Qy        301 GGGACACGACTGGACATTCAA 321
              ||||||||||||||| ||| ||
Db        301 GGGACACGACTGGAGATTAAA 321


Search completed: September 12, 2005, 02:01:34
Job time : 510.25 secs
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